

# DECLARATION AND POWER OF ATTORNEY FOR PATENT APPLICATION

As the below named inventor, I hereby declare that:

My residence, post office address and citizenship is as stated below next to my name;

I believe that I am the original, first and sole inventor (if only one name is listed below) or an original, first and joint inventor (if plural names are listed below) of the subject matter which is claimed and for which a patent is sought on the invention entitled:

## NUCLEIC ACIDS AND PROTEINS FROM GROUP B STREPTOCOCCUS

the specification of which: ☒ is attached hereto.  
☐ was filed on: \_\_\_\_\_  
as Application No.: \_\_\_\_\_  
and was amended on: \_\_\_\_\_ (if applicable).

I have reviewed and understand the contents of the above-identified specification, including the claims, as amended by any amendment referred to above. I acknowledge the duty to disclose information which is material to patentability as defined in 37 C.F.R. § 1.56.

And I hereby authorize and request our agents, Brobeck, Phleger & Harrison LLP, whose address is set forth below, to insert above, the filing date and application number of said application when known.

### Prior Foreign Application(s)

I hereby claim foreign priority benefits under Title 35, United States Code, § 119(a)-(d) or § 365(b) of any foreign application(s) for patent or inventor's certificate, or § 365(a) of any PCT international application which designated at least one country other than the United States of America, listed below and have also identified below any foreign application(s) for patent or inventor's certificate having a filing date before that of the application on which priority is claimed:

Country	Application Number	Date of Filing (day, month, year)	Date of Issue (day, month, year)	Priority Claimed	
German	9816335.5	July 27, 1998		Yes <input checked="" type="checkbox"/>	No <input type="checkbox"/>
				Yes <input type="checkbox"/>	No <input type="checkbox"/>

### Prior Provisional Application(s)

I hereby claim the benefit under Title 35, United States Code § 119(e) of any United States provisional application(s) listed below:

Application Number	Date of Filing (day, month, year)
60/125,163	March 19, 1999

### Prior United States Application(s)

I hereby claim the benefit under Title 35, United States Code, § 120 of any United States application(s), or § 365(c) of any PCT international application designating the United States of America, listed below and, insofar as the subject matter of each of the claims of this application is not disclosed in the prior United States application in the manner provided by the first paragraph of Title 35, United States Code, § 112, I acknowledge the duty to disclose material information as defined in Title 37, Code of Federal Regulations, § 1.56(a) which occurred between the filing date of the prior application and the national or PCT international filing date of this application:

Application Number	Date of Filing (day, month, year)	Status - Patented, Pending, Abandoned
PCT/GB99/02444	27 July 1999	Pending

And I hereby appoint, both jointly and severally, as my attorneys with full power of substitution and revocation, to prosecute this application and to transact all business in the Patent and Trademark Office connected herewith the following attorneys, their registration numbers being listed after their names:

Rodger L. Tate, Registration No. 27,399; Anthony W. Shaw, Registration No. 30,104; James Remenick, Registration No. 36,902; Michael J. Songer, Reg. No. 39,841; Cono A. Carrano, Registration No. 39,623; Laurence H. Posorske, Registration No. 34,698; Floyd B. Chapman, Registration No. 40,555; David J. Kulik, Registration No. 36,576; Robert A. King, Registration No. 42,738; and Trevor Q. Coddington, Registration No. 46,633.

All correspondence and telephone communications should be addressed to: Brobeck, Phleger & Harrison LLP; Intellectual Property Department; 1333 H Street, N.W.; Suite 800; Washington, DC 20005; telephone number (202) 220-6000; facsimile number (202) 220-5200, which is also the address, telephone and facsimile numbers of each of the above listed attorneys.

I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine and imprisonment, or both, under 18 U.S.C. § 1001, and that such willful false statements may jeopardize the validity of the application or any patent issuing thereon.

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Signature \_\_\_\_\_ Date \_\_\_\_\_

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Citizenship: Ireland

Residence: Cambridge, United Kingdom

Post Office  
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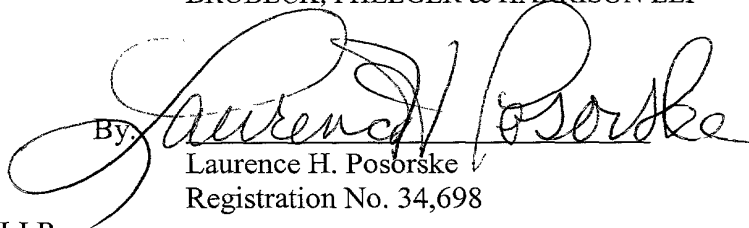
It is believed that no fees are required for this submission; however, the Commissioner is authorized to charge any fee necessary for entry of this paper to Deposit Account 50-1640.

Respectfully submitted,

BROBECK, PHLEGER & HARRISON LLP

January 26, 2001

By

  
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## SEQUENCE LISTING

<110> Microbial Technics Limited

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Hanniffy, Sean B

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<120> Proteins

<130> PWC/P21122WO

<140> PCT/GB99/02452

<141> 1999-07-27

<150> GB 9816336.3

<151> 1998-07-27

<150> US 60/125329

<151> 1999-03-19

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<213> Streptococcus pneumoniae

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20

25

30

Gly Ile Ser Val Gly Ile Gly His Leu Gln Gly Ser Ser Met Ala Lys

35

40

45

Asn Asn Lys Val Ala Val Val Thr Thr Val Pro Ser Val Ala Glu Gly

50

55

60

Leu Lys Asn Val Asn Gly Val Asn Phe Asp Tyr Lys Asp Glu Ala Ser

65

70

75

80

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Lys Arg Leu Ala Gln Thr Ile Gln Phe Thr Glu Lys Ile Asp Glu Ala  
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Lys Glu Asn Lys Lys Phe Ile Gln Thr Ile Ala Ala Gly Ala Leu Gly  
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Phe Phe Leu Tyr Met Ile Leu Ile Thr Tyr Ala Gly Val Thr Ala Gln  
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Glu Val Ala Ser Glu Lys Gly Thr Lys Ile Met Glu Val Val Phe Ser  
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Ser Ile Arg Ala Ser His Tyr Phe Tyr Ala Arg Met Met Ala Leu Phe  
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Gly Asp Asn Leu Leu Leu Lys Ile Gly Ser Tyr Ile Pro Phe Ile Ser  
 325 330 335

Thr Phe Phe Met Pro Phe Arg Thr Ile Asn Asp Tyr Ala Gly Gly Ala  
 340 345 350

Glu Ala Trp Ile Ser Leu Ala Ile Thr Val Ile Phe Ala Val Val Ala  
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<213> Streptococcus pneumoniae

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<212> PRT

<213> Streptococcus pneumoniae

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Val Ile Ser Ala Tyr Asn Glu Glu Lys Tyr Leu Pro Gly Leu Ile Glu
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Asp Leu Lys Asn Gln Thr Tyr Pro Lys Glu Asp Ile Glu Ile Leu Phe
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Ile Lys Glu Asp Thr Glu Phe Asn Ser Ile Arg Leu Tyr Asn Asn Pro
      85             90             95

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Lys Lys Asn Gln Ala Ser Gly Phe Asn Leu Gly Val Lys His Ser Val
      100            105            110

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 145 150 155 160

Thr Leu His Leu Val Glu Glu Asn Met Phe Gly Ser Ser Ile Ala Asn  
 165 170 175

Tyr Arg Asn Ser Ser Glu Asp Arg Tyr Val Ser Ser Ile Phe His Gly  
 180 185 190

Met Tyr Lys Arg Glu Val Phe Gln Lys Val Gly Leu Val Asn Glu Gln  
 195 200 205

Leu Gly Arg Thr Glu Asp Asn Asp Ile His Tyr Arg Ile Arg Glu Tyr  
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Gly Tyr Lys Ile Arg Tyr Ser Pro Ser Ile Leu Ser Tyr Gln Tyr Ile  
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Arg Pro Thr Phe Lys Lys Met Leu His Gln Lys Tyr Ser Asn Gly Leu  
 245 250 255

Trp Ile Gly Leu Thr Ser His Val Gln Phe Lys Cys Leu Ser Leu Phe  
 260 265 270

His Tyr Val Pro Cys Leu Phe Val Leu Ser Leu Val Phe Ser Leu Ala  
 275 280 285

Leu Leu Pro Ile Thr Phe Val Phe Ile Thr Leu Leu Leu Gly Ala Tyr  
 290 295 300

Phe Leu Leu Leu Ser Leu Leu Thr Leu Leu Thr Leu Leu Lys His Lys  
 305 310 315 320

Asn Gly Phe Leu Ile Val Met Pro Phe Ile Leu Phe Ser Ile His Phe  
 325 330 335

Ala Tyr Gly Leu Gly Thr Ile Val Gly Leu Ile Arg Gly Phe Lys Trp  
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<212> DNA

<213> Streptococcus pneumoniae

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<211> 231

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<213> Streptococcus pneumoniae

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Thr Gly Ala Gly Ala Phe Ala Tyr Ser Thr Phe Ile Val Lys Pro Glu  
 35 40 45

Tyr Thr Ser Thr Thr Arg Ile Tyr Val Val Asn Arg Asn Gln Gly Asp  
 50 55 60

Lys Pro Gly Leu Thr Asn Gln Asp Leu Gln Ala Gly Thr Tyr Leu Val  
 65 70 75 80

Lys Asp Tyr Arg Glu Ile Ile Leu Ser Gln Asp Val Leu Glu Glu Val  
 85 90 95

Val Ser Asp Leu Lys Leu Asp Leu Thr Pro Lys Gly Leu Ala Asn Lys  
 100 105 110

Ile Lys Val Thr Val Pro Val Asp Thr Arg Ile Val Ser Ile Ser Val  
 115 120 125

Asn Asp Arg Val Pro Glu Glu Ala Ser Arg Ile Ala Asn Ser Leu Arg  
 130 135 140

Glu Val Ala Ala Gln Lys Ile Ile Ser Ile Thr Arg Val Ser Asp Val  
 145 150 155 160

Thr Thr Leu Glu Glu Ala Arg Pro Ala Ile Ser Pro Ser Ser Pro Asn  
 165 170 175

Ile Lys Arg Asn Thr Leu Ile Gly Phe Leu Ala Gly Val Ile Gly Thr  
 180 185 190

Ser Val Ile Val Leu His Leu Glu Leu Leu Asp Thr Arg Val Lys Arg  
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<213> Streptococcus pneumoniae

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Arg Ala Asp His Val Phe Asp Gly Asp Leu Ser Asp Tyr Asp Met Ile  
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Val Leu Pro Gly Gly Met Pro Gly Ser Ala His Leu Arg Asp Asn Gln  
65 70 75 80

Thr Leu Ile Gln Glu Leu Gln Ser Phe Glu Gln Glu Gly Lys Lys Leu  
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Ala Ala Ile Cys Ala Ala Pro Ile Ala Leu Asn Gln Ala Glu Ile Leu  
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Lys Asn Lys Arg Tyr Thr Cys Tyr Asp Gly Val Gln Glu Gln Ile Leu  
115 120 125

Asp Gly His Tyr Val Lys Glu Thr Val Val Val Asp Gly Gln Leu Thr  
130 135 140

Thr Ser Arg Gly Pro Ser Thr Ala Leu Ala Phe Ala Tyr Glu Leu Val  
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Arg Asp Val Phe Gly Lys Asn Gln  
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&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 9

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&lt;210&gt; 10

&lt;211&gt; 101

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 10

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      20              25             30

Leu Glu Thr Thr Lys Lys Leu Pro Thr Thr Asn Glu Gln Leu Gln Ala
      35              40             45

Val Arg Leu Ser Gly Leu Val Asn Arg Glu Leu Leu Leu Asn Pro Lys
      50              55             60

His Pro Ala Pro Glu Leu Leu Asn Leu Ala Arg Phe Val Lys Arg Glu
      65              70             75             80

Glu Ala Lys Tyr Arg Gly Thr Ala Thr Ser Ala Leu Met Tyr Glu Glu
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Leu Phe Lys Met Leu

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<212> DNA

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<213> Streptococcus pneumoniae

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Leu Leu Phe Val Gly Ile Gln Ser Asp Gly Ile Lys Ser Leu Leu Ser  
 35 40 45

Met Ser Lys Glu Pro Val Tyr Asp Ser Arg Thr Glu Lys Leu Thr Phe  
 50 55 60

Gly Lys Glu Val Glu Asn Leu Glu Ile Thr Leu His Gln His Thr Leu  
 65 70 75 80

Thr Ile Thr Asp Ser Phe Asp Asp Gln Ile His Ile Ser Tyr His Pro  
 85 90 95

Ser Leu Ser Ala His His Asp Leu Ile Thr Asn Gln Asn Asp Arg Thr  
 100 105 110

Leu Ser Leu Thr Asp Lys Lys Leu Ser Glu Thr Pro Phe Leu Ser Ser  
 115 120 125

Gly Ile Gly Gly Ile Leu His Ile Ala Ser Ser Tyr Ser Ser Arg Phe  
 130 135 140

Glu Glu Val Ile Leu Arg Leu Pro Lys Gly Arg Thr Leu Lys Gly Ile  
 145 150 155 160

Asn Ile Ser Ala Asn Arg Gly Gln Thr Thr Ile Ile Asn Ala Ser Leu  
 165 170 175

Glu Asn Ala Thr Leu Asn Thr Asn Ser Tyr Ile Leu Arg Ile Glu Gly  
 180 185 190

Ser Arg Ile Lys Asn Ser Lys Leu Thr Thr Pro Asn Ile Val Asn Ile  
 195 200 205

Phe Asp Thr Val Leu Thr Asp Ser Gln Leu Glu Ser Thr Glu Asn His  
 210 215 220

Phe His Ala Glu Asn Ile Gln Val His Gly Lys Val Glu Leu Thr Ala  
 225 230 235 240

Lys Asp Tyr Leu Arg Ile Ile Leu Asp Gln Lys Glu Ser Gln Arg Ile  
 245 250 255

Asn Trp Asp Ile Ser Ser Asn Tyr Gly Ser Ile Phe Gln Phe Thr Arg  
 260 265 270

Glu Lys Pro Glu Ser Arg Gly Thr Glu Leu Ser Asn Pro Tyr Lys Thr  
 275 280 285

Glu Lys Thr Asp Val Lys Asp Gln Leu Ile Ala Arg Ser Asp Asp Asn  
 290 295 300

Ile Asp Leu Ile Ser Thr Pro Ser Arg Arg  
 305 310

<210> 13

<211> 879

<212> DNA

<213> Streptococcus pneumoniae

<400> 13

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 ccaattgaaa aaaataactca gttagaggag gaagtccttc aagctgaagt cgaattggaa 180  
 agccagcaag aagagaaaaat tgaagctcct gaagacagtg aagcgagaac agaaatagaa 240  
 gaaaagaagg catctaatto tactgaagaa gagccagacc tttctaaaga aacagaaaaa 300  
 gtcactatag ctgaagagag ccaagaagct cttcctcagc aaaaagcaac cacgaaagag 360  
 ccacttctta tcagtaaatac tttagaaagt ccttatatcc ccgaccaagc tccaaaatct 420  
 agggataaat ggaaagagca agtgcttgat ttttggtctt ggctagtgga agcgatcaaa 480  
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 attctgtttt ctgcatcttc ctttttcttt agtatctatc acatcaaaca tgcttactat 600  
 ggacatatag caagcattaa cagtcgcttc cctgagcagc tagctccttt aactcttttt 660  
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ttcgttgatga gacgatttat ccaccaggaa aaggactgga cgctagacaa ggttctccaa 780  
 caatatagtc aactcttggc aattccaatc tctcactgc tattgctagt ttctttgctt 840  
 tctttgatag cctacgattt acagccctct tgttgtgtga 879

<210> 14

<211> 292

<212> PRT

<213> Streptococcus pneumoniae

<400> 14

Met Lys Gln Glu Trp Phe Glu Ser Asn Asp Phe Val Lys Thr Thr Ser  
 1 5 10 15

Lys Asn Lys Pro Glu Glu Gln Ala Gln Glu Val Ala Asp Lys Ala Glu  
 20 25 30

Glu Thr Ile Ala Asp Leu Asp Thr Pro Ile Glu Lys Asn Thr Gln Leu  
 35 40 45

Glu Glu Glu Val Pro Gln Ala Glu Val Glu Leu Glu Ser Gln Gln Glu  
 50 55 60

Glu Lys Ile Glu Ala Pro Glu Asp Ser Glu Ala Arg Thr Glu Ile Glu  
 65 70 75 80

Glu Lys Lys Ala Ser Asn Ser Thr Glu Glu Glu Pro Asp Leu Ser Lys  
 85 90 95

Glu Thr Glu Lys Val Thr Ile Ala Glu Glu Ser Gln Glu Ala Leu Pro  
 100 105 110

Gln Gln Lys Ala Thr Thr Lys Glu Pro Leu Leu Ile Ser Lys Ser Leu  
 115 120 125

Glu Ser Pro Tyr Ile Pro Asp Gln Ala Pro Lys Ser Arg Asp Lys Trp  
 130 135 140

Lys Glu Gln Val Leu Asp Phe Trp Ser Trp Leu Val Glu Ala Ile Lys  
 145 150 155 160

Ser Pro Thr Ser Lys Leu Glu Thr Ser Ile Thr His Ser Tyr Thr Ala  
 165 170 175

Phe Leu Leu Leu Ile Leu Phe Ser Ala Ser Ser Phe Phe Phe Ser Ile  
 180 185 190

Tyr His Ile Lys His Ala Tyr Tyr Gly His Ile Ala Ser Ile Asn Ser  
 195 200 205

Arg Phe Pro Glu Gln Leu Ala Pro Leu Thr Leu Phe Ser Ile Ile Ser  
 210 215 220

Ile Leu Val Ala Thr Thr Leu Phe Phe Phe Ser Phe Leu Leu Gly Ser  
 225 230 235 240

Phe Val Val Arg Arg Phe Ile His Gln Glu Lys Asp Trp Thr Leu Asp  
 245 250 255

Lys Val Leu Gln Gln Tyr Ser Gln Leu Leu Ala Ile Pro Ile Ser Ser  
 260 265 270

Leu Leu Leu Leu Val Ser Leu Leu Ser Leu Ile Ala Tyr Asp Leu Gln  
 275 280 285

Pro Ser Cys Val  
 290

<210> 15

<211> 990

<212> DNA

<213> Streptococcus pneumoniae

&lt;400&gt; 15

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 actaccgtta tcggctttat cctgcttttt gtaggtatcc aatctgacgg gattaagagc 180  
 ctactttcca tgtccaaaga acctgtctat gatagccgta cggaaaagct aacctttggc 240  
 aaggaagtcg aaaacctaga aattactctc caccaacaca cgctcaccat cacagactct 300  
 ttcgatgata aaatccacat ttcttaccat ccatctcttt ctgctcacca tgatcttata 360  
 accaatcaga acgatagaac tctgagtctc actgataaga aactgtctga aactccggtt 420  
 ctctcttctg gaattggtgg gattcttcat atcgcaagta gctactctag tcgttttgaa 480  
 gaagttattc tccgactacc aaaagggaga actctaaaag ggatcaacat ctgagccaat 540  
 cgcggaacaa ccaccatcat aaatgctagc cttgaaaatg cgaccctcaa taaaaacagc 600  
 tatactctcc gaattgaagg aagtcgtatc aaaaacagta aactcacaac gcccaatata 660  
 gttaatatct ttgatacagt tcttacagat agtcagctag agtcaacaga gaatcacttc 720  
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 atcatcctag accagaaaga aagccaacga attaactggg acatctcaag caactatggt 840  
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&lt;210&gt; 16

&lt;211&gt; 329

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 16

Met Gln Leu Ala Ser Ser Val Tyr Ser Leu Phe Val Trp Tyr Asn Leu

1

5

10

15

Phe Leu Lys Lys Glu Arg Glu Val Ile Ser Met Arg Lys Trp Thr Lys

20

25

30

Gly Phe Leu Ile Phe Gly Val Val Thr Thr Val Ile Gly Phe Ile Leu

35

40

45

Leu Phe Val Gly Ile Gln Ser Asp Gly Ile Lys Ser Leu Leu Ser Met

50

55

60

Ser Lys Glu Pro Val Tyr Asp Ser Arg Thr Glu Lys Leu Thr Phe Gly  
65 70 75 80

Lys Glu Val Glu Asn Leu Glu Ile Thr Leu His Gln His Thr Leu Thr  
85 90 95

Ile Thr Asp Ser Phe Asp Asp Gln Ile His Ile Ser Tyr His Pro Ser  
100 105 110

Leu Ser Ala His His Asp Leu Ile Thr Asn Gln Asn Asp Arg Thr Leu  
115 120 125

Ser Leu Thr Asp Lys Lys Leu Ser Glu Thr Pro Phe Leu Ser Ser Gly  
130 135 140

Ile Gly Gly Ile Leu His Ile Ala Ser Ser Tyr Ser Ser Arg Phe Glu  
145 150 155 160

Glu Val Ile Leu Arg Leu Pro Lys Gly Arg Thr Leu Lys Gly Ile Asn  
165 170 175

Ile Ser Ala Asn Arg Gly Gln Thr Thr Ile Ile Asn Ala Ser Leu Glu  
180 185 190

Asn Ala Thr Leu Asn Thr Asn Ser Tyr Ile Leu Arg Ile Glu Gly Ser  
195 200 205

Arg Ile Lys Asn Ser Lys Leu Thr Thr Pro Asn Ile Val Asn Ile Phe  
210 215 220

Asp Thr Val Leu Thr Asp Ser Gln Leu Glu Ser Thr Glu Asn His Phe  
225 230 235 240

His Ala Glu Asn Ile Gln Val His Gly Lys Val Glu Leu Thr Ala Lys  
245 250 255

Asp Tyr Leu Arg Ile Ile Leu Asp Gln Lys Glu Ser Gln Arg Ile Asn  
260 265 270

Trp Asp Ile Ser Ser Asn Tyr Gly Ser Ile Phe Gln Phe Thr Arg Glu  
 275 280 285

Lys Pro Glu Ser Arg Gly Thr Glu Leu Ser Asn Pro Tyr Lys Thr Glu  
 290 295 300

Lys Thr Asp Val Lys Asp Gln Leu Ile Ala Arg Ser Asp Asp Asn Ile  
 305 310 315 320

Asp Leu Ile Ser Thr Pro Ser Arg Arg  
 325

<210> 17

<211> 79

<212> DNA

<213> Streptococcus pneumoniae

<400> 17

atgatatgta aaatgaagca gggagggagc agggcgctgct ggggatggag agtgggggag 60  
 ggacgctgct attttaatc 79

<210> 18

<211> 26

<212> PRT

<213> Streptococcus pneumoniae

<400> 18

Met Ile Cys Lys Met Lys Gln Gly Gly Ser Arg Ala Cys Trp Gly Trp  
 1 5 10 15

Arg Val Gly Glu Gly Arg Cys Tyr Phe Asn  
 20 25



&lt;210&gt; 19

&lt;211&gt; 715

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 19

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cgataaaagag gccttgagta atctcaatth gcagattgaa aatggagaga ttatgggctt 60
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accagcaggt ggtcgtatth ttgtagacgg tcaggagtta tcggaaaatc gcttggctat 180
taaacgaaag attggctacg tagcagactc gcctgactta tttttacgct taacggccaa 240
tgaattttgg gaattgatcg cctcatccta tgatctgagt agatctgact tggaggctag 300
tctagctagg ctattgaacg tttttgattt tgctgaaaat cgctatcagg ttattgaaac 360
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agaggtggca gagcaagtct gtgatcggat tgccattttg aaaaaggggc atttgattta 600
ttgtggtaag gtagaggact tgaggaaaga ccaccagac cagtctttgg aaagtatcta 660
ccttagtctt gctggtagaa aagaggaggt tgcggatgcg tctcaaggtc attaa 715

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&lt;210&gt; 20

&lt;211&gt; 237

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 20

Asp Lys Glu Ala Leu Ser Asn Leu Asn Leu Gln Ile Glu Asn Gly Glu

1

5

10

15

Ile Met Gly Leu Ile Gly His Asn Gly Ala Gly Lys Ser Thr Thr Ile

20

25

30

Lys Ser Leu Val Ser Ile Ile Ser Pro Ser Ser Gly Arg Ile Leu Val

35

40

45

Asp Gly Gln Glu Leu Ser Glu Asn Arg Leu Ala Ile Lys Arg Lys Ile

50

55

60

Gly Tyr Val Ala Asp Ser Pro Asp Leu Phe Leu Arg Leu Thr Ala Asn  
 65 70 75 80

Glu Phe Trp Glu Leu Ile Ala Ser Ser Tyr Asp Leu Ser Arg Ser Asp  
 85 90 95

Leu Glu Ala Ser Leu Ala Arg Leu Leu Asn Val Phe Asp Phe Ala Glu  
 100 105 110

Asn Arg Tyr Gln Val Ile Glu Thr Leu Ser His Gly Met Arg Gln Lys  
 115 120 125

Val Phe Val Ile Gly Ala Leu Leu Ser Asp Pro Asp Ile Trp Val Leu  
 130 135 140

Asp Glu Pro Leu Thr Gly Leu Asp Pro Gln Ala Ala Phe Asp Leu Lys  
 145 150 155 160

Gln Met Met Lys Glu His Ala Gln Lys Gly Lys Thr Val Leu Phe Ser  
 165 170 175

Thr His Val Leu Glu Val Ala Glu Gln Val Cys Asp Arg Ile Ala Ile  
 180 185 190

Leu Lys Lys Gly His Leu Ile Tyr Cys Gly Lys Val Glu Asp Leu Arg  
 195 200 205

Lys Asp His Pro Asp Gln Ser Leu Glu Ser Ile Tyr Leu Ser Leu Ala  
 210 215 220

Gly Arg Lys Glu Glu Val Ala Asp Ala Ser Gln Gly His  
 225 230 235

<210> 21

<211> 360

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 21

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tgttgccagt tttcagaaaag aatttttagca acttggctaa agaaactact gctagtctct 180
tcagttgttg tagcttcggc aggttggtcc ttgatcatat gatccatcaa ggcaacttgg 240
tcattctttg aaatggtttc aatgctggca ttgatttggc taatacgatt gtcattttta 300
cgaagcccgga tagcgatagc tgtatcttct tccccagttt tgaaaccagg ttctacttga 360

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<210> 22

<211> 119

<212> PRT

<213> *Streptococcus pneumoniae*

<400> 22

Met Ala Leu Phe Ser Glu Arg Gly Ala Val Arg Lys Thr Pro Met Ala

1

5

10

15

Ser Pro Ile Met Arg Pro Met Met Val Pro Thr Ile Glu Ile Lys Arg

20

25

30

Val Ile Pro Ala Pro Arg Lys Ser Cys Cys Gln Phe Ser Glu Arg Ile

35

40

45

Leu Ala Thr Trp Leu Lys Lys Leu Leu Leu Val Ser Ser Val Val Val

50

55

60

Ala Ser Ala Gly Cys Ser Leu Ile Ile Arg Ser Ile Lys Ala Thr Trp

65

70

75

80

Ser Ser Phe Glu Met Val Ser Met Leu Ala Leu Ile Trp Leu Ile Arg

85

90

95

Leu Ser Phe Leu Arg Ser Pro Ile Ala Ile Ala Val Ser Ser Ser Pro  
 100 105 110

Val Leu Lys Pro Gly Ser Thr  
 115

<210> 23

<211> 1455

<212> DNA

<213> Streptococcus pneumoniae

<400> 23

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 gcagctcatg ctgataatgt tcgaactaaa gatgaaatca atcgtcaaaa acaagaacat 480  
 gtcaaagata atgagaaggt taactctaata gttgctgtag caagggtctca gggacgatat 540  
 acgacaaatg atgggttatgt ctttaatcca gctgatatta tcgaagatac gggtaatgct 600  
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 gaattagcag cagctaaagc acatctggct ggaaaaaata tgcaaccgag tcagttaagc 720  
 tattcttcaa cagctagtga caataacacg caatctgtag caaaaggatc aactagcaag 780  
 ccagcaaata aatctgaaaa tctccagagt cttttgaagg aactctatga ttcacctagc 840  
 gcccaacgtt acagtgaatc agatggcctg gtctttgacc ctgctaagat tatcagtcgt 900  
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 cattacattc caaatcaaaa tcaaattggg caaccgactc ttccaaacaa tagtctagca 1260  
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cgcaaaaaca tttag

1455

&lt;210&gt; 24

&lt;211&gt; 484

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 24

Met Lys Phe Ser Lys Lys Tyr Ile Ala Ala Gly Ser Ala Val Ile Val

1

5

10

15

Ser Leu Ser Leu Cys Ala Tyr Ala Leu Asn Gln His Arg Ser Gln Glu

20

25

30

Asn Lys Asp Asn Asn Arg Val Ser Tyr Val Asp Gly Ser Gln Ser Ser

35

40

45

Gln Lys Ser Glu Asn Leu Thr Pro Asp Gln Val Ser Gln Lys Glu Gly

50

55

60

Ile Gln Ala Glu Gln Ile Val Ile Lys Ile Thr Asp Gln Gly Tyr Val

65

70

75

80

Thr Ser His Gly Asp His Tyr His Tyr Tyr Asn Gly Lys Val Pro Tyr

85

90

95

Asp Ala Leu Phe Ser Glu Glu Leu Leu Met Lys Asp Pro Asn Tyr Gln

100

105

110

Leu Lys Asp Ala Asp Ile Val Asn Glu Val Lys Gly Gly Tyr Ile Ile

115

120

125

Lys Val Asp Gly Lys Tyr Tyr Val Tyr Leu Lys Asp Ala Ala His Ala

130

135

140

Asp Asn Val Arg Thr Lys Asp Glu Ile Asn Arg Gln Lys Gln Glu His

145

150

155

160

Val Lys Asp Asn Glu Lys Val Asn Ser Asn Val Ala Val Ala Arg Ser  
 165 170 175

Gln Gly Arg Tyr Thr Thr Asn Asp Gly Tyr Val Phe Asn Pro Ala Asp  
 180 185 190

Ile Ile Glu Asp Thr Gly Asn Ala Tyr Ile Val Pro His Gly Gly His  
 195 200 205

Tyr His Tyr Ile Pro Lys Ser Asp Leu Ser Ala Ser Glu Leu Ala Ala  
 210 215 220

Ala Lys Ala His Leu Ala Gly Lys Asn Met Gln Pro Ser Gln Leu Ser  
 225 230 235 240

Tyr Ser Ser Thr Ala Ser Asp Asn Asn Thr Gln Ser Val Ala Lys Gly  
 245 250 255

Ser Thr Ser Lys Pro Ala Asn Lys Ser Glu Asn Leu Gln Ser Leu Leu  
 260 265 270

Lys Glu Leu Tyr Asp Ser Pro Ser Ala Gln Arg Tyr Ser Glu Ser Asp  
 275 280 285

Gly Leu Val Phe Asp Pro Ala Lys Ile Ile Ser Arg Thr Pro Asn Gly  
 290 295 300

Val Ala Ile Pro His Gly Asp His Tyr His Phe Ile Pro Tyr Ser Lys  
 305 310 315 320

Leu Ser Ala Leu Glu Glu Lys Ile Ala Arg Met Val Pro Ile Ser Gly  
 325 330 335

Thr Gly Ser Thr Val Ser Thr Asn Ala Lys Pro Asn Glu Val Val Ser  
 340 345 350

Ser Leu Gly Ser Leu Ser Ser Asn Pro Ser Ser Leu Thr Thr Ser Lys  
 355 360 365

Glu Leu Ser Ser Ala Ser Asp Gly Tyr Ile Phe Asn Pro Lys Asp Ile  
 370 375 380

Val Glu Glu Thr Ala Thr Ala Tyr Ile Val Arg His Gly Asp His Phe  
 385 390 395 400

His Tyr Ile Pro Lys Ser Asn Gln Ile Gly Gln Pro Thr Leu Pro Asn  
 405 410 415

Asn Ser Leu Ala Thr Pro Ser Pro Ser Leu Pro Ile Asn Pro Gly Thr  
 420 425 430

Ser His Glu Lys His Glu Glu Asp Gly Tyr Gly Phe Asp Ala Asn Arg  
 435 440 445

Ile Ile Ala Glu Asp Glu Ser Gly Phe Val Met Ser His Gly Asp His  
 450 455 460

Asn His Tyr Phe Phe Lys Lys Asp Leu Thr Glu Glu Gln Ile Lys Val  
 465 470 475 480

Arg Lys Asn Ile

<210> 25

<211> 840

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 25

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 gaagttcagt tgggagtacc taaaattacg gccttttgta aacgattggc tgatagaggc 780  
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<210> 26

<211> 279

<212> PRT

<213> Streptococcus pneumoniae

<400> 26

Met Gly Ile Ala Leu Glu Asn Val Asn Phe Thr Tyr Gln Glu Gly Thr

1

5

10

15

Pro Leu Ala Ser Ala Ala Leu Ser Asp Val Ser Leu Thr Ile Glu Asp

20

25

30

Gly Ser Tyr Thr Ala Leu Ile Gly His Thr Gly Ser Gly Lys Ser Thr

35

40

45

Ile Leu Gln Leu Leu Asn Gly Leu Leu Val Pro Ser Gln Gly Ser Val

50

55

60

Arg Val Phe Asp Thr Leu Ile Thr Ser Thr Ser Lys Asn Lys Asp Ile

65

70

75

80

Arg Gln Ile Arg Lys Gln Val Gly Leu Val Phe Gln Phe Ala Glu Asn

85

90

95

Gln Ile Phe Glu Glu Thr Val Leu Lys Asp Val Ala Phe Gly Pro Gln

100

105

110



Asn Phe Gly Val Ser Glu Glu Asp Ala Val Lys Thr Ala Arg Glu Lys  
 115 120 125

Leu Ala Leu Val Gly Ile Asp Glu Ser Leu Phe Asp Arg Ser Pro Phe  
 130 135 140

Glu Leu Ser Gly Gly Gln Met Arg Arg Val Ala Ile Ala Gly Ile Leu  
 145 150 155 160

Ala Met Glu Pro Ala Ile Leu Val Leu Asp Glu Pro Thr Ala Gly Leu  
 165 170 175

Asp Pro Leu Gly Arg Lys Glu Leu Met Thr Leu Phe Lys Lys Leu His  
 180 185 190

Gln Ser Gly Met Thr Ile Val Leu Val Thr His Leu Met Asp Asp Val  
 195 200 205

Ala Glu Tyr Ala Asn Gln Val Tyr Val Met Glu Lys Gly Arg Leu Val  
 210 215 220

Lys Gly Gly Lys Pro Ser Asp Val Phe Gln Asp Val Val Phe Met Glu  
 225 230 235 240

Glu Val Gln Leu Gly Val Pro Lys Ile Thr Ala Phe Cys Lys Arg Leu  
 245 250 255

Ala Asp Arg Gly Val Ser Phe Lys Arg Leu Pro Ile Lys Ile Glu Glu  
 260 265 270

Phe Lys Glu Ser Leu Asn Gly  
 275

&lt;210&gt; 27

&lt;211&gt; 6360

&lt;212&gt; DNA

<213> *Streptococcus pneumoniae*

&lt;400&gt; 27

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cctgtgatag acaataacac tagcaatgaa gaagcaaaaa tcaaagaaga aaattccaat 240  
aaatcccaag gagattatac ggactcattt gtgaataaaa acacagaaaa tccccaaaaa 300  
gaagataaag ttgtctatat tgctgaattt aaagataaag aatctggaga aaaagcaatc 360  
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ggcaaaatca ctgtagaaaa atatgatgat ggaagggatt attttgaccc acatgggatg 840  
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&lt;210&gt; 28

&lt;211&gt; 2119

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 28

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Ile His Ser Ala Met Glu Thr Ser Gln Asp Phe Lys Glu Lys Lys Thr  
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Ala Val Ile Lys Glu Lys Glu Val Val Ser Lys Asn Pro Val Ile Asp  
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Asn Asn Thr Ser Asn Glu Glu Ala Lys Ile Lys Glu Glu Asn Ser Asn  
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Asn Pro Lys Lys Glu Asp Lys Val Val Tyr Ile Ala Glu Phe Lys Asp  
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Thr Lys Val Leu Tyr Thr Tyr Asp Arg Ile Phe Asn Gly Ser Ala Ile  
 130 135 140

Glu Thr Thr Pro Asp Asn Leu Asp Lys Ile Lys Gln Ile Glu Gly Ile  
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Ser Ser Val Glu Arg Ala Gln Lys Val Gln Pro Met Met Asn His Ala  
 165 170 175

Lys Ala Gly Ile Pro Met Val Val Ala Thr Gly Asn Tyr Ala Thr Ser  
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Ala Ser Ser Ser Ser Trp Asp Leu Val Ala Asn Asn His Leu Lys Met  
 385 390 395 400

Thr Asp Thr Gly Asn Val Thr Arg Thr Ala Ala His Glu Asp Ala Ile  
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Ala Val Ala Ser Ala Lys Asn Gln Thr Val Glu Phe Asp Lys Val Asn  
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Ile Gly Gly Glu Ser Phe Lys Tyr Arg Asn Ile Gly Ala Phe Phe Asp  
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Lys Ser Lys Ile Thr Thr Asn Glu Asp Gly Thr Lys Ala Pro Ser Lys  
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Asp Leu Lys Asn Ala Phe Lys Lys Ala Met Asp Lys Gly Ala Arg Ala  
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Glu Leu Pro Ala Met Gly Tyr Glu Ala Asp Glu Gly Thr Lys Ser Gln  
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Asn Pro Asp Lys Lys Thr Glu Val Lys Arg Asn Asn Lys Glu Asp Phe  
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Lys Asp Lys Leu Glu Gln Tyr Tyr Pro Ile Asp Met Glu Ser Phe Asn  
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Asp	Val	Ser	Ala	Pro	Gly	Lys	Asn	Ile	Lys	Ser	Thr	Leu	Asn	Val	Ile
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Pro	Ile	Val	Ala	Ala	Ser	Thr	Val	Leu	Ile	Arg	Pro	Lys	Leu	Lys	Glu
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Met	Leu	Glu	Arg	Pro	Val	Leu	Lys	Asn	Leu	Lys	Gly	Asp	Asp	Lys	Ile
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Pro	Arg	Gln	Gln	Gly	Ala	Gly	Leu	Ile	Asn	Val	Ala	Asn	Ala	Leu	Arg
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Asn	Ser	Tyr	Gly	Ser	Ile	Ser	Leu	Lys	Glu	Ile	Lys	Gly	Asp	Lys	Lys
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Tyr	Phe	Thr	Ile	Lys	Leu	His	Asn	Thr	Ser	Asn	Arg	Pro	Leu	Thr	Phe
785					790						795				800



Phe Asn Asn Glu Gly Ile Asn Ala Pro Ser Ser Ser Gly Ser Lys Ile  
995 1000 1005

Ala Asn Ile Tyr Pro Leu Asp Ser Asn Gly Asn Pro Gln Asp Ala Gln  
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Leu Glu Arg Gly Leu Thr Pro Ser Pro Leu Val Leu Arg Ser Ala Glu  
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Glu Gly Leu Ile Ser Ile Val Asn Thr Asn Lys Glu Gly Glu Asn Gln  
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Arg Asp Leu Lys Val Ile Ser Arg Glu His Phe Ile Arg Gly Ile Leu  
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Asn Ser Lys Ser Asn Asp Ala Lys Gly Ile Lys Ser Ser Lys Leu Lys  
 1075 1080 1085

Val Trp Gly Asp Leu Lys Trp Asp Gly Leu Ile Tyr Asn Pro Arg Gly  
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Arg Glu Glu Asn Ala Pro Glu Ser Lys Asp Asn Gln Asp Pro Ala Thr  
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Lys Phe Lys Tyr Arg Leu Thr Lys Asp Tyr Pro Trp Gln Val Ser Tyr  
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Ile Pro Val Lys Ile Asp Asn Thr Ala Pro Lys Ile Val Ser Val Asp  
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Phe Ser Asn Pro Glu Lys Ile Lys Leu Ile Thr Lys Asp Thr Tyr His  
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Lys Val Lys Asp Gln Tyr Lys Asn Glu Thr Leu Phe Ala Arg Asp Gln  
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Lys Glu His Pro Glu Lys Phe Asp Glu Ile Ala Asn Glu Val Trp Tyr  
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Glu Val Thr Tyr Ala Gly Glu Gly Gln Gly Arg Asn Arg Lys Leu Asp  
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Gly Lys Ile Ile Glu Val Ile Ala Leu Asp Gly Ser Ser Asn Phe Thr  
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Lys Ile His Arg Ile Lys Phe Ala Asn Gln Ala Asp Glu Lys Gly Met  
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Ile Ser Tyr Tyr Leu Val Asp Pro Asp Gln Asp Ser Ser Lys Tyr Gln  
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Lys Leu Gly Glu Ile Ala Glu Ser Lys Phe Lys Asn Leu Gly Asn Gly  
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Lys Glu Gly Ser Leu Lys Lys Asp Thr Thr Gly Val Glu His His His  
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Gln Glu Asn Glu Glu Ser Ile Lys Glu Lys Ser Ser Phe Thr Ile Asp  
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Arg Asn Ile Ser Thr Ile Arg Asp Phe Glu Asn Lys Asp Leu Lys Lys  
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Leu Ile Lys Lys Lys Phe Arg Glu Val Asp Asp Phe Thr Ser Glu Thr  
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Gly Lys Arg Met Glu Glu Tyr Asp Tyr Lys Tyr Asp Asp Lys Gly Asn  
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Ile Ile Ala Tyr Asp Asp Gly Thr Asp Leu Glu Tyr Glu Thr Glu Lys  
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Leu Asp Glu Ile Lys Ser Lys Ile Tyr Gly Val Leu Ser Pro Ser Lys  
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Asp Gly His Phe Glu Ile Leu Gly Lys Ile Ser Asn Val Ser Lys Asn  
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Ala Lys Val Tyr Tyr Gly Asn Asn Tyr Lys Ser Ile Glu Ile Lys Ala  
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Ser Tyr Gly Asn Val Ile Glu Leu Gly Glu Gly Asp Leu Ser Lys Asn  
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Lys Pro Asp Asn Leu Thr Lys Met Glu Ser Gly Lys Ile Tyr Ser Asp  
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Ser Glu Lys Gln Gln Tyr Leu Leu Lys Asp Asn Ile Ile Leu Arg Lys  
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Gly Tyr Ala Leu Lys Val Thr Thr Tyr Asn Pro Gly Lys Thr Asp Met  
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Asp Ser Arg Asn Val Glu Asp Gly Arg Ser Thr Gln Ser Val Leu Met  
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Ser Ala Leu Asp Gly Phe Asn Ile Ile Arg Tyr Gln Val Phe Thr Phe  
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Lys Met Asn Asp Lys Gly Glu Ala Ile Asp Lys Asp Gly Asn Leu Val  
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Thr Asp Ser Ser Lys Leu Val Leu Phe Gly Lys Asp Asp Lys Glu Tyr  
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Thr Gly Glu Asp Lys Phe Asn Val Glu Ala Ile Lys Glu Asp Gly Ser  
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Met Leu Phe Ile Asp Thr Lys Pro Val Asn Leu Ser Met Asp Lys Asn  
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Tyr Phe Asn Pro Ser Lys Ser Asn Lys Ile Tyr Val Arg Asn Pro Glu  
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Phe Tyr Leu Arg Gly Lys Ile Ser Asp Lys Gly Gly Phe Asn Trp Glu  
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Leu Arg Val Asn Glu Ser Val Val Asp Asn Tyr Leu Ile Tyr Gly Asp  
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Leu His Ile Asp Asn Thr Arg Asp Phe Asn Ile Lys Leu Asn Val Lys  
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Asp Gly Asp Ile Met Asp Trp Gly Met Lys Asp Tyr Lys Ala Asn Gly  
 1795 1800 1805

Phe Pro Asp Lys Val Thr Asp Met Asp Gly Asn Val Tyr Leu Gln Thr  
 1810 1815 1820

Gly Tyr Ser Asp Leu Asn Ala Lys Ala Val Gly Val His Tyr Gln Phe  
 1825 1830 1835 1840

Leu Tyr Asp Asn Val Lys Pro Glu Val Asn Ile Asp Pro Lys Gly Asn  
 1845 1850 1855

Thr Ser Ile Glu Tyr Ala Asp Gly Lys Ser Val Val Phe Asn Ile Asn  
 1860 1865 1870

Asp Lys Arg Asn Asn Gly Phe Asp Gly Glu Ile Gln Glu Gln His Ile  
 1875 1880 1885

Tyr Ile Asn Gly Lys Glu Tyr Thr Ser Phe Asn Asp Ile Lys Gln Ile  
 1890 1895 1900

Ile Asp Lys Thr Leu Asn Ile Lys Ile Val Val Lys Asp Phe Ala Arg  
 1905 1910 1915 1920

Asn Thr Thr Val Lys Glu Phe Ile Leu Asn Lys Asp Thr Gly Glu Val  
 1925 1930 1935

Ser Glu Leu Lys Pro His Arg Val Thr Val Thr Ile Gln Asn Gly Lys  
 1940 1945 1950

Glu Met Ser Ser Thr Ile Val Ser Glu Glu Asp Phe Ile Leu Pro Val  
 1955 1960 1965

Tyr Lys Gly Glu Leu Glu Lys Gly Tyr Gln Phe Asp Gly Trp Glu Ile  
 1970 1975 1980

Ser Gly Phe Glu Gly Lys Lys Asp Ala Gly Tyr Val Ile Asn Leu Ser  
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Lys Asp Thr Phe Ile Lys Pro Val Phe Lys Lys Ile Glu Glu Lys Lys  
 2005 2010 2015

Glu Glu Glu Asn Lys Pro Thr Phe Asp Val Ser Lys Lys Lys Asp Asn  
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Pro Gln Val Asn His Ser Gln Leu Asn Glu Ser His Arg Lys Glu Asp  
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Leu Gln Arg Glu Glu His Ser Gln Lys Ser Asp Ser Thr Lys Asp Val  
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Thr Ala Thr Val Leu Asp Lys Asn Asn Ile Ser Ser Lys Ser Thr Thr  
 2065 2070 2075 2080

Asn Asn Pro Asn Lys Leu Pro Lys Thr Gly Thr Ala Ser Gly Ala Gln  
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<212> DNA

<213> Streptococcus pneumoniae

<400> 29

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&lt;211&gt; 198

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 30

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Ala Pro Ser Ala His Asn Ser Gln Pro Trp Lys Phe Val Val Val Arg  
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Glu Lys Asn Ala Glu Leu Ala Lys Leu Ala Tyr Gly Ser Asn Phe Glu  
 50 55 60

Gln Val Ser Ser Ala Pro Val Thr Ile Ala Leu Phe Thr Asp Thr Asp  
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Leu Ala Lys Arg Ala Arg Lys Ile Ala Arg Val Gly Gly Ala Asn Asn  
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Phe Ser Glu Glu Gln Leu Gln Tyr Phe Met Lys Asn Leu Pro Ala Glu  
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Phe Ala Arg Tyr Ser Glu Gln Gln Val Ser Asp Tyr Leu Ala Leu Asn  
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Ala Gly Leu Val Ala Met Asn Leu Val Leu Ala Leu Thr Asp Gln Gly  
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Ile Gly Ser Asn Ile Ile Leu Gly Phe Asp Lys Ser Lys Val Asn Glu  
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Val Leu Glu Ile Glu Asp Arg Phe Arg Pro Glu Leu Leu Ile Thr Val  
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Gly Tyr Thr Asp Glu Lys Leu Glu Pro Ser Tyr Arg Leu Pro Val Asp  
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Glu Ile Ile Glu Lys Arg  
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<211> 1401

<212> DNA

<213> Streptococcus pneumoniae

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 acttaccttg ccctcttct cagccagttt ggctttgctg gtccagccaa agactacctt 900  
 gacatcgag gtaaaattct cttgaacgat catgagggtg aaaatcttaa gattgctcat 960  
 gtggatgaaa agatgggtgc tctttctatg aatgccggcg tcttccactt cgatgaaaca 1020  
 agtgctgata ataccattgc cctcaacatc cgctatccaa aaggaacaag tccagaacaa 1080  
 atcaagtcaa tccttgaaaa cttgccagtt gtttctgtta gcctgtctga acacggtcac 1140  
 acgcctcact atgtgccaat ggaagatcca cttgtgcaaa cttgttgaa tatctatgaa 1200  
 aaacaaactg gctttaaagg tcatgaacaa gtcacggtg gtggaacctt tggctgcttg 1260  
 ctagaacgcg gagttgccta cgggtgctatg ttcccagact cgattgatac catgcaccaa 1320  
 gccaatgaat ttatgcctt ggatgatctt ttccgagcag cagcaattta tgccgaagct 1380  
 atttacgaat tgatcaaata a 1401

&lt;210&gt; 32

&lt;211&gt; 466

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 32

Met Thr Ala Ile Asp Phe Thr Ala Glu Val Glu Lys Arg Lys Glu Asp

1

5

10

15

Leu Leu Ala Asp Leu Phe Ser Leu Leu Glu Ile Asn Ser Glu Arg Asp

20

25

30

Asp Ser Lys Ala Asp Ala Gln His Pro Phe Gly Pro Gly Pro Val Lys

35

40

45

Ala Leu Glu Lys Phe Leu Glu Ile Ala Asp Arg Asp Gly Tyr Pro Thr

50

55

60

Lys Asn Val Asp Asn Tyr Ala Gly His Phe Glu Phe Gly Asp Gly Glu

65

70

75

80

Glu Val Leu Gly Ile Phe Ala His Met Asp Val Val Pro Ala Gly Ser

85

90

95

Gly Trp Asp Thr Asp Pro Tyr Thr Pro Thr Ile Lys Asp Gly Arg Leu

100

105

110

Tyr Ala Arg Gly Ala Ser Asp Asp Lys Gly Pro Thr Thr Ala Cys Tyr

115

120

125

Tyr Gly Leu Lys Ile Ile Lys Glu Leu Gly Leu Pro Thr Ser Lys Lys

130

135

140

Val Arg Phe Ile Val Gly Thr Asp Glu Glu Ser Gly Trp Ala Asp Met

145

150

155

160

Pro Lys Gly Thr Ser Pro Glu Gln Ile Lys Ser Ile Leu Glu Asn Leu  
355 360 365

Pro Val Val Ser Val Ser Leu Ser Glu His Gly His Thr Pro His Tyr  
 370 375 380

Val Pro Met Glu Asp Pro Leu Val Gln Thr Leu Leu Asn Ile Tyr Glu  
 385 390 395 400

Lys Gln Thr Gly Phe Lys Gly His Glu Gln Val Ile Gly Gly Gly Thr  
 405 410 415

Phe Gly Arg Leu Leu Glu Arg Gly Val Ala Tyr Gly Ala Met Phe Pro  
 420 425 430

Asp Ser Ile Asp Thr Met His Gln Ala Asn Glu Phe Ile Ala Leu Asp  
 435 440 445

Asp Leu Phe Arg Ala Ala Ala Ile Tyr Ala Glu Ala Ile Tyr Glu Leu  
 450 455 460

Ile Lys  
 465

<210> 33

<211> 1617

<212> DNA

<213> Streptococcus pneumoniae

<400> 33

gtgtatacta ttataaaatc aaatataaaa aaatttagtt tattaacgat atttattggt 60  
 gctggtcaat tattgctaatt ttatgcagca actattaatg ctctggtggt gaatgaatta 120  
 attgcatga atttagagcg gtttttgaaa ttgtcaatct accaaatgat tgtctggtgt 180  
 gggataatat tccttgactg ggtagtgaat aattatcagg ttgaagtgat ccaagagttt 240  
 aatctagaga ttcgaaatag agttgccaca gacatctcta actctaccta tcaagaattt 300  
 catagtaaatt catcaggaac atatctttcg tggctaaata atgatgttca gactttaaat 360  
 gatcaggcgt ttaaacaact ttttttagta ataaaaggaa tttctggtac tatatttgca 420  
 gttgtgactc ttaatcacta tcattggtca ttgactgtag ccaccttggt ttcattaatg 480  
 attatgctac ttgtaccaa aatctttgca tcgaaaatgc gagaagttag tctaaattta 540

actaaccacaaa atgaagcttt tttaaaatct agtgagacta tattgaatgg atttgatgtg 600  
 ttagcgtcct tgaatctttt atatgtattg cctaagaaaa ttaaagaagc aggaatttta 660  
 ttaaagatgg ttatacaaag aaagacaact gtagaaacgt tagcaggcgc tattagcttc 720  
 tttctcaata ttttttttca gatatctctc gtttttttaa caggctatct tgcaataaaa 780  
 ggaatagtga aaattggtac tattgaagca ataggagcac taacaggtgt tattttttaca 840  
 gcgctaggtg aattaggagg tcaattatcc tctattattg gtacgaagcc tattttttta 900  
 aaattgtatt caattaatcc aattgagtca aataaaatga atgatatoga accaaatgag 960  
 gtgaatagag attttccggt atatgaagca aaaaatattt gctataagta tggagataaa 1020  
 gaaatattaa aaaacttaaa tttttgtttt caacgtaatg aaaagtattt aatttttaggt 1080  
 gaaagtggaa gcgggaaatc tacattatta aaattattga atggcttttt gagagattat 1140  
 agtggagaat tgcgattctg cggggatgat ataaaaaaaa cctcctattt aaatatgggt 1200  
 tcgaatgttc tatatgtaga tcaaaaagct tttttgtttg aaggtaacgat tagagataat 1260  
 attttattgg aagaaaatta tactgatgaa gaaatactac agtcttttaga gcaagttggt 1320  
 ttgagtgtaa aagattttcc taataacatt ttagattatt atgttggtga tgatgggaga 1380  
 ttactgtcag gagggcagaa acaaaaaaatt acttttagcta gagggctaata tagaaataag 1440  
 aaaatagtat taattgacga gggaaacttct gctatcgata ggagaacttc gttagcgatt 1500  
 gaacgtaaga tattagatag agaggatttg actgtcatta ttgttaccba tgctccgcatt 1560  
 ccggaactta aacaatattt tactaagata tatcaatttc caaaggattt tatttaa 1617

<210> 34

<211> 538

<212> PRT

<213> Streptococcus pneumoniae

<400> 34

Met Tyr Thr Ile Ile Lys Ser Asn Ile Lys Lys Phe Ser Leu Leu Thr

1

5

10

15

Ile Phe Ile Val Ala Gly Gln Leu Leu Leu Ile Tyr Ala Ala Thr Ile

20

25

30

Asn Ala Leu Val Leu Asn Glu Leu Ile Ala Met Asn Leu Glu Arg Phe

35

40

45

Leu Lys Leu Ser Ile Tyr Gln Met Ile Val Trp Cys Gly Ile Ile Phe

50

55

60

Leu Asp Trp Val Val Lys Asn Tyr Gln Val Glu Val Ile Gln Glu Phe  
65 70 75 80

Asn Leu Glu Ile Arg Asn Arg Val Ala Thr Asp Ile Ser Asn Ser Thr  
85 90 95

Tyr Gln Glu Phe His Ser Lys Ser Ser Gly Thr Tyr Leu Ser Trp Leu  
100 105 110

Asn Asn Asp Val Gln Thr Leu Asn Asp Gln Ala Phe Lys Gln Leu Phe  
115 120 125

Leu Val Ile Lys Gly Ile Ser Gly Thr Ile Phe Ala Val Val Thr Leu  
130 135 140

Asn His Tyr His Trp Ser Leu Thr Val Ala Thr Leu Phe Ser Leu Met  
145 150 155 160

Ile Met Leu Leu Val Pro Lys Ile Phe Ala Ser Lys Met Arg Glu Val  
165 170 175

Ser Leu Asn Leu Thr Asn Gln Asn Glu Ala Phe Leu Lys Ser Ser Glu  
180 185 190

Thr Ile Leu Asn Gly Phe Asp Val Leu Ala Ser Leu Asn Leu Leu Tyr  
195 200 205

Val Leu Pro Lys Lys Ile Lys Glu Ala Gly Ile Leu Leu Lys Met Val  
210 215 220

Ile Gln Arg Lys Thr Thr Val Glu Thr Leu Ala Gly Ala Ile Ser Phe  
225 230 235 240

Phe Leu Asn Ile Phe Phe Gln Ile Ser Leu Val Phe Leu Thr Gly Tyr  
245 250 255

Leu Ala Ile Lys Gly Ile Val Lys Ile Gly Thr Ile Glu Ala Ile Gly  
260 265 270

Ala Leu Thr Gly Val Ile Phe Thr Ala Leu Gly Glu Leu Gly Gly Gln  
 275 280 285

Leu Ser Ser Ile Ile Gly Thr Lys Pro Ile Phe Leu Lys Leu Tyr Ser  
 290 295 300

Ile Asn Pro Ile Glu Ser Asn Lys Met Asn Asp Ile Glu Pro Asn Glu  
 305 310 315 320

Val Asn Arg Asp Phe Pro Leu Tyr Glu Ala Lys Asn Ile Cys Tyr Lys  
 325 330 335

Tyr Gly Asp Lys Glu Ile Leu Lys Asn Leu Asn Phe Cys Phe Gln Arg  
 340 345 350

Asn Glu Lys Tyr Leu Ile Leu Gly Glu Ser Gly Ser Gly Lys Ser Thr  
 355 360 365

Leu Leu Lys Leu Leu Asn Gly Phe Leu Arg Asp Tyr Ser Gly Glu Leu  
 370 375 380

Arg Phe Cys Gly Asp Asp Ile Lys Lys Thr Ser Tyr Leu Asn Met Val  
 385 390 395 400

Ser Asn Val Leu Tyr Val Asp Gln Lys Ala Tyr Leu Phe Glu Gly Thr  
 405 410 415

Ile Arg Asp Asn Ile Leu Leu Glu Glu Asn Tyr Thr Asp Glu Glu Ile  
 420 425 430

Leu Gln Ser Leu Glu Gln Val Gly Leu Ser Val Lys Asp Phe Pro Asn  
 435 440 445

Asn Ile Leu Asp Tyr Tyr Val Gly Asp Asp Gly Arg Leu Leu Ser Gly  
 450 455 460

Gly Gln Lys Gln Lys Ile Thr Leu Ala Arg Gly Leu Ile Arg Asn Lys  
 465 470 475 480

Lys Ile Val Leu Ile Asp Glu Gly Thr Ser Ala Ile Asp Arg Arg Thr  
 485 490 495

Ser Leu Ala Ile Glu Arg Lys Ile Leu Asp Arg Glu Asp Leu Thr Val  
 500 505 510

Ile Ile Val Thr His Ala Pro His Pro Glu Leu Lys Gln Tyr Phe Thr  
 515 520 525

Lys Ile Tyr Gln Phe Pro Lys Asp Phe Ile  
 530 535

<210> 35

<211> 705

<212> DNA

<213> Streptococcus pneumoniae

<400> 35

ataacagtta aacagattat ggacgaaata gccgtttcag atatgactgc aaggcgctat 60  
 ttacaggaat tagctgataa agatttgctg attcgtgtgc atggtggagc tgaaaaactt 120  
 cgaaccaact cccttttgac taatgagcga tcaaatattg aaaaacaagc cctccaaacg 180  
 gcagaaaaac aagaaatagc ccattttgca ggcagtctag tagaagaaag agaaactatt 240  
 ttcattggac caggaacaac attagagttt tttgcgcgtg agttgcctat tgacaatatc 300  
 cgcgtcgtaa ccaacagtct acctgttttt ctgattttta gcgaacgaaa attaacagat 360  
 ttgattttta taggtggaaa ttatcgcgat attacaggtg cttttgttgg tacattgacc 420  
 ctacaaaatc tctctaactt ccaattttct aaagctttcg ttagctgtaa tggatttcaa 480  
 aacggagctc tagctacttt tagcgaggaa gagggagagg ctcaacgcat cgctttaaat 540  
 aattctaata aaaaatattt actgcagat catagcaagt tcaataagtt tgatttttat 600  
 actttttata atgtatcaaa tcttgatact attgtttcag attctaaaact aagtgattca 660  
 atccttttta agctatctaa acacattaaa gtcacgaagc cttaa 705

<210> 36

<211> 234

<212> PRT

<213> Streptococcus pneumoniae



&lt;400&gt; 36

Ile Thr Val Lys Gln Ile Met Asp Glu Ile Ala Val Ser Asp Met Thr  
 1 5 10 15

Ala Arg Arg Tyr Leu Gln Glu Leu Ala Asp Lys Asp Leu Leu Ile Arg  
 20 25 30

Val His Gly Gly Ala Glu Lys Leu Arg Thr Asn Ser Leu Leu Thr Asn  
 35 40 45

Glu Arg Ser Asn Ile Glu Lys Gln Ala Leu Gln Thr Ala Glu Lys Gln  
 50 55 60

Glu Ile Ala His Phe Ala Gly Ser Leu Val Glu Glu Arg Glu Thr Ile  
 65 70 75 80

Phe Ile Gly Pro Gly Thr Thr Leu Glu Phe Phe Ala Arg Glu Leu Pro  
 85 90 95

Ile Asp Asn Ile Arg Val Val Thr Asn Ser Leu Pro Val Phe Leu Ile  
 100 105 110

Leu Ser Glu Arg Lys Leu Thr Asp Leu Ile Leu Ile Gly Gly Asn Tyr  
 115 120 125

Arg Asp Ile Thr Gly Ala Phe Val Gly Thr Leu Thr Leu Gln Asn Leu  
 130 135 140

Ser Asn Leu Gln Phe Ser Lys Ala Phe Val Ser Cys Asn Gly Ile Gln  
 145 150 155 160

Asn Gly Ala Leu Ala Thr Phe Ser Glu Glu Glu Gly Glu Ala Gln Arg  
 165 170 175

Ile Ala Leu Asn Asn Ser Asn Lys Lys Tyr Leu Leu Ala Asp His Ser  
 180 185 190

Lys Phe Asn Lys Phe Asp Phe Tyr Thr Phe Tyr Asn Val Ser Asn Leu  
 195 200 205

Asp Thr Ile Val Ser Asp Ser Lys Leu Ser Asp Ser Ile Leu Phe Lys  
 210 215 220

Leu Ser Lys His Ile Lys Val Ile Lys Pro  
 225 230

<210> 37

<211> 483

<212> DNA

<213> Streptococcus pneumoniae

<400> 37

atgactgagt ttctgtaga tcttcttcta gaagccatta aactagctcg ttggacctac 60  
 tactatcact tgaaacagct agacaaaaca gataaagacc aagagcttaa aactgaaatt 120  
 caatccatct ttatcgaaca caagggaat tatgcttata gccgggttca tttagaacta 180  
 agaaatcggtg gttatctggt aaatcataaa agagttcaag gcttgatgaa agtactcaat 240  
 ttacaagcta aaatgcgaaa gaaacgaaaa tattcttctc ataaaggaga cgttggtaag 300  
 aaggcagaga atctcattca agcccaattt gaaggctcta aaacaatgga aaagtgtac 360  
 acagatgtga ctgaatttgc cattccagca agtactcaaa agctttactt atcaccagtt 420  
 ttagatggct ttaacagcga aattattgct tttaatcttt cttgttcgcc taatttagaa 480  
 taa 483

<210> 38

<211> 160

<212> PRT

<213> Streptococcus pneumoniae

<400> 38

Met Thr Glu Phe Ser Leu Asp Leu Leu Leu Glu Ala Ile Lys Leu Ala  
 1 5 10 15

Arg Trp Thr Tyr Tyr Tyr His Leu Lys Gln Leu Asp Lys Thr Asp Lys  
 20 25 30

Asp Gln Glu Leu Lys Thr Glu Ile Gln Ser Ile Phe Ile Glu His Lys  
 35 40 45

Gly Asn Tyr Ala Tyr Arg Arg Val His Leu Glu Leu Arg Asn Arg Gly  
 50 55 60

Tyr Leu Val Asn His Lys Arg Val Gln Gly Leu Met Lys Val Leu Asn  
 65 70 75 80

Leu Gln Ala Lys Met Arg Lys Lys Arg Lys Tyr Ser Ser His Lys Gly  
 85 90 95

Asp Val Gly Lys Lys Ala Glu Asn Leu Ile Gln Ala Gln Phe Glu Gly  
 100 105 110

Ser Lys Thr Met Glu Lys Cys Tyr Thr Asp Val Thr Glu Phe Ala Ile  
 115 120 125

Pro Ala Ser Thr Gln Lys Leu Tyr Leu Ser Pro Val Leu Asp Gly Phe  
 130 135 140

Asn Ser Glu Ile Ile Ala Phe Asn Leu Ser Cys Ser Pro Asn Leu Glu  
 145 150 155 160

<210> 39

<211> 1266

<212> DNA

<213> Streptococcus pneumoniae

<400> 39

ccaggatttg gtaccgttgc aagtgggtgtg cctttcctcc taaaggaaaa tggaggaaaa 60  
 atcaatcaat cagcacattc agatatcaaa gttgctaagg tattggtcaa ggatgaagat 120  
 gaaaaaaatc gcttgcttgc agcagggaat gactttaact ttgtaaccaa tgtggatgat 180

attttatcag accaggatat tactatcgta gtggaattga tggggcgat tgagcctgct 240  
 aaaaccttta tcaactcgtag cttggaagct ggaaaacacg ttgttactgc taacaaggac 300  
 ctttttagctg tccatggcgc agaattgcta gaaatcgctc aagctaacaa ggtagcactt 360  
 tactacgaag cagcagttgc tgggtgggatt ccaattcttc gtacttttagc aaattccttg 420  
 gcttctgata aaattacgag cgtgcttgga gtagtcaacg gaacttccaa cttcatgggtg 480  
 accaagatgg tggagaagg ctggtcttac gatgatgctc ttgcggaagc acaacgtcta 540  
 ggattttgcag aaagcgatcc gacgaatgac gtagatggga ttgatgcagc ctacaagatg 600  
 gttattttga gccaatgtgc ctttggcatg aagattgctt ttgatgatgt agcccacaag 660  
 ggaatccgca atatcacacc agaagacgta gctgtagctc aagagcttgg ttacgtagtg 720  
 aaattgggtg gttctattga ggaaacttct tcaggtattg ctgcagaagt gactccaacc 780  
 ttcctaccta aagcgacccc acttgctagt gtgaatggcg taatgaacgc tgtctttgta 840  
 gaatctatcg gtattggtga gtctatgtac tacggaccag gtgcgggtca aaaaccaact 900  
 gcaacaagtg ttgtagctga tattgtccgt atcgttcgtc gtttgaatga tggtagctatt 960  
 ggcaaagact tcaacgaata tagccgtgac ttggtcttgg caaatcctga agatgtcaaa 1020  
 gcaaactact atttctcaat cttggctcta gactcaaaaag gtcaggtctt gaagttggct 1080  
 gaaatcttca atgctcaaga tatttctctt aagcaaatcc ttcaagatgg caaagagggt 1140  
 gacaaggcgc gtgtcggtat catcacacac aagattaata aagcccagct tgaaaatgtc 1200  
 tcagctgaat tgaagaaggt ttcagaattc gacctcttga ataccttcaa ggtgctagga 1260  
 gaataa 1266

<210> 40

<211> 421

<212> PRT

<213> Streptococcus pneumoniae

<400> 40

Pro Gly Phe Gly Thr Val Ala Ser Gly Val Pro Phe Leu Leu Lys Glu

1

5

10

15

Asn Gly Gly Lys Ile Asn Gln Ser Ala His Ser Asp Ile Lys Val Ala

20

25

30

Lys Val Leu Val Lys Asp Glu Asp Glu Lys Asn Arg Leu Leu Ala Ala

35

40

45

Gly Asn Asp Phe Asn Phe Val Thr Asn Val Asp Asp Ile Leu Ser Asp

50

55

60

Gln Asp Ile Thr Ile Val Val Glu Leu Met Gly Arg Ile Glu Pro Ala  
65 70 75 80

Lys Thr Phe Ile Thr Arg Ala Leu Glu Ala Gly Lys His Val Val Thr  
85 90 95

Ala Asn Lys Asp Leu Leu Ala Val His Gly Ala Glu Leu Leu Glu Ile  
100 105 110

Ala Gln Ala Asn Lys Val Ala Leu Tyr Tyr Glu Ala Ala Val Ala Gly  
115 120 125

Gly Ile Pro Ile Leu Arg Thr Leu Ala Asn Ser Leu Ala Ser Asp Lys  
130 135 140

Ile Thr Arg Val Leu Gly Val Val Asn Gly Thr Ser Asn Phe Met Val  
145 150 155 160

Thr Lys Met Val Glu Glu Gly Trp Ser Tyr Asp Asp Ala Leu Ala Glu  
165 170 175

Ala Gln Arg Leu Gly Phe Ala Glu Ser Asp Pro Thr Asn Asp Val Asp  
180 185 190

Gly Ile Asp Ala Ala Tyr Lys Met Val Ile Leu Ser Gln Phe Ala Phe  
195 200 205

Gly Met Lys Ile Ala Phe Asp Asp Val Ala His Lys Gly Ile Arg Asn  
210 215 220

Ile Thr Pro Glu Asp Val Ala Val Ala Gln Glu Leu Gly Tyr Val Val  
225 230 235 240

Lys Leu Val Gly Ser Ile Glu Glu Thr Ser Ser Gly Ile Ala Ala Glu  
245 250 255

Val Thr Pro Thr Phe Leu Pro Lys Ala His Pro Leu Ala Ser Val Asn  
260 265 270

Gly Val Met Asn Ala Val Phe Val Glu Ser Ile Gly Ile Gly Glu Ser  
 275 280 285

Met Tyr Tyr Gly Pro Gly Ala Gly Gln Lys Pro Thr Ala Thr Ser Val  
 290 295 300

Val Ala Asp Ile Val Arg Ile Val Arg Arg Leu Asn Asp Gly Thr Ile  
 305 310 315 320

Gly Lys Asp Phe Asn Glu Tyr Ser Arg Asp Leu Val Leu Ala Asn Pro  
 325 330 335

Glu Asp Val Lys Ala Asn Tyr Tyr Phe Ser Ile Leu Ala Leu Asp Ser  
 340 345 350

Lys Gly Gln Val Leu Lys Leu Ala Glu Ile Phe Asn Ala Gln Asp Ile  
 355 360 365

Ser Phe Lys Gln Ile Leu Gln Asp Gly Lys Glu Gly Asp Lys Ala Arg  
 370 375 380

Val Val Ile Ile Thr His Lys Ile Asn Lys Ala Gln Leu Glu Asn Val  
 385 390 395 400

Ser Ala Glu Leu Lys Lys Val Ser Glu Phe Asp Leu Leu Asn Thr Phe  
 405 410 415

Lys Val Leu Gly Glu  
 420

<210> 41

<211> 1725

<212> DNA

<213> Streptococcus pneumoniae

<400> 41

atgaaacacc tattatctta cttcaaacc tacatcaagg aatcaatttt agcccccttg 60  
 ttcaagctgt tagaagctgt ttttgagctc ttggttccca tgggtgattgc tgggattggt 120

gaccaatctt tacctcaggg agatcaaggt catctctgga tgcagattgg cctgctcctt 180  
 atctttgcag taattggcgt tttagtggcc ttgatagctc aattttactc agcaaaggca 240  
 gcagtaggtt ctgctaagga attgacaaac gatctttatc gtcataattct ttccttgccc 300  
 aaggacagca gagaccgtct gacaacttct agtttggtca ctgccttgac ttcggatacc 360  
 taccagattc agactggtat caatcaattc ctgcgtctct ttttacgagc gccattatc 420  
 gtttttggtg ccatttttat ggcttatcga atctcagctg agttgacttt ctgggtctta 480  
 gtcttggttg ccattttgac cattgtcatt gtagggttat ctcgattggg caatcctttc 540  
 tacagtagtc tcagaaagaa aacggaccaa ctgggtcagg aaacgcgcca gcaattgcaa 600  
 gggatgcggg ttattcgtgc ttttggtcaa gaaaaacgag agttacagat ttttcaaacc 660  
 cttaaccaag tttatgctag attacaagaa aagacaggtt tctgggtctag tttattaaca 720  
 cctctgacct atctgattgt caatggaact cttctcgtaa ttatctggca aggctatat 780  
 tcaattcaag gaggagtgtc cagtcaaggt gctctcattg ctcttatcaa ttacctctta 840  
 cagattttgg tggaattggg caagctagcc atgttgatca attccctcaa ccagtcctat 900  
 atctcagtca agcgaatcga ggaagtcttt gttgaggctc cagaggatat ccattcagag 960  
 ttagaacaaa agcaagctac cagagataag gttttacaag tccaagaatt gacctttacc 1020  
 tatcctgatg cggcccagcc ttctctgaga tacatttctt ttgatatgac tcaaggacaa 1080  
 attctaggtt tcatcggggg aactggttct ggtaaataca gcttggtgca actcttactt 1140  
 ggactttatc cagtagacaa ggggaacatt gacctttatc aaaatggacg tagtcctctt 1200  
 aatttgagc agtggcggtc ttggattgcc tatgtacctc aaaaggctga actctttaa 1260  
 ggaaccattc gttccaactt gactctaggt ttcaatcaag aagtatctga ccaggaactc 1320  
 tggcaggcct tggagattgc gcaagctaag gattttgtca gtgaaaagga aggactcttg 1380  
 gatgctctag ttgaggcagg ggggcgaaat ttctcaggtg gacaaaaaca aagattgtct 1440  
 atcgcccag cagtcttgcg ccaggctccg tttctcatcc tagatgatgc aacctcggca 1500  
 ctggatacca ttacagagtc caagctcttg aaagctatta gagaaaattt tccaaacacg 1560  
 agcttaattt tgatctctca acgaacctca actttacaga tggcggacca gattctcctc 1620  
 ttggaaaaag gtgagttgct agctgttggc aagcacgatg acttgatgaa atccagccaa 1680  
 gtctattgtg aaatcaatgc atccaacat ggaaaggagg actag 1725

<210> 42

<211> 574

<212> PRT

<213> *Streptococcus pneumoniae*

<400> 42

Met Lys His Leu Leu Ser Tyr Phe Lys Pro Tyr Ile Lys Glu Ser Ile

1

5

10

15

Leu Ala Pro Leu Phe Lys Leu Leu Glu Ala Val Phe Glu Leu Leu Val  
 20 25 30

Pro Met Val Ile Ala Gly Ile Val Asp Gln Ser Leu Pro Gln Gly Asp  
 35 40 45

Gln Gly His Leu Trp Met Gln Ile Gly Leu Leu Leu Ile Phe Ala Val  
 50 55 60

Ile Gly Val Leu Val Ala Leu Ile Ala Gln Phe Tyr Ser Ala Lys Ala  
 65 70 75 80

Ala Val Gly Ser Ala Lys Glu Leu Thr Asn Asp Leu Tyr Arg His Ile  
 85 90 95

Leu Ser Leu Pro Lys Asp Ser Arg Asp Arg Leu Thr Thr Ser Ser Leu  
 100 105 110

Val Thr Arg Leu Thr Ser Asp Thr Tyr Gln Ile Gln Thr Gly Ile Asn  
 115 120 125

Gln Phe Leu Arg Leu Phe Leu Arg Ala Pro Ile Ile Val Phe Gly Ala  
 130 135 140

Ile Phe Met Ala Tyr Arg Ile Ser Ala Glu Leu Thr Phe Trp Phe Leu  
 145 150 155 160

Val Leu Val Ala Ile Leu Thr Ile Val Ile Val Gly Leu Ser Arg Leu  
 165 170 175

Val Asn Pro Phe Tyr Ser Ser Leu Arg Lys Lys Thr Asp Gln Leu Val  
 180 185 190

Gln Glu Thr Arg Gln Gln Leu Gln Gly Met Arg Val Ile Arg Ala Phe  
 195 200 205

Gly Gln Glu Lys Arg Glu Leu Gln Ile Phe Gln Thr Leu Asn Gln Val  
 210 215 220



Glu Leu Phe Lys Gly Thr Ile Arg Ser Asn Leu Thr Leu Gly Phe Asn  
420 425 430

Gln Glu Val Ser Asp Gln Glu Leu Trp Gln Ala Leu Glu Ile Ala Gln  
 435 440 445

Ala Lys Asp Phe Val Ser Glu Lys Glu Gly Leu Leu Asp Ala Leu Val  
 450 455 460

Glu Ala Gly Gly Arg Asn Phe Ser Gly Gly Gln Lys Gln Arg Leu Ser  
 465 470 475 480

Ile Ala Arg Ala Val Leu Arg Gln Ala Pro Phe Leu Ile Leu Asp Asp  
 485 490 495

Ala Thr Ser Ala Leu Asp Thr Ile Thr Glu Ser Lys Leu Leu Lys Ala  
 500 505 510

Ile Arg Glu Asn Phe Pro Asn Thr Ser Leu Ile Leu Ile Ser Gln Arg  
 515 520 525

Thr Ser Thr Leu Gln Met Ala Asp Gln Ile Leu Leu Leu Glu Lys Gly  
 530 535 540

Glu Leu Leu Ala Val Gly Lys His Asp Asp Leu Met Lys Ser Ser Gln  
 545 550 555 560

Val Tyr Cys Glu Ile Asn Ala Ser Gln His Gly Lys Glu Asp  
 565 570

<210> 43

<211> 1224

<212> DNA

<213> Streptococcus pneumoniae

<400> 43

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<210> 44
<211> 407
<212> PRT
<213> Streptococcus pneumoniae
```

Met Lys Arg Ser Leu Asp Ser Arg Val Asp Tyr Ser Leu Leu Leu Pro

Val Phe Phe Leu Leu Val Ile Gly Val Val Ala Ile Tyr Ile Ala Val

Ser His Asp Tyr Pro Asn Asn Ile Leu Pro Ile Leu Gly Gln Gln Val

Ala Trp Ile Ala Leu Gly Leu Val Ile Gly Phe Val Val Met Leu Phe

50                      55                      60

Gln Ile Ala Ile Gly Ser Gly Gly Leu Phe Gly Gln Gly Phe Asn Ala  
260 265 270

Ser Asn Leu Leu Ile Pro Val Arg Glu Ser Asp Met Ile Phe Thr Val  
 275 280 285

Ile Ala Glu Asp Phe Gly Phe Ile Gly Ser Val Leu Val Ile Ala Leu  
 290 295 300

Tyr Leu Met Leu Ile Tyr Arg Met Leu Lys Ile Thr Leu Lys Ser Asn  
 305 310 315 320

Asn Gln Phe Tyr Thr Tyr Ile Ser Thr Gly Leu Ile Met Met Leu Leu  
 325 330 335

Phe His Ile Phe Glu Asn Ile Gly Ala Val Thr Gly Leu Leu Pro Leu  
 340 345 350

Thr Gly Ile Pro Leu Pro Phe Ile Ser Gln Gly Gly Ser Ala Ile Ile  
 355 360 365

Ser Asn Leu Ile Gly Val Gly Leu Leu Leu Ser Met Ser Tyr Gln Thr  
 370 375 380

Asn Leu Ala Glu Glu Lys Ser Gly Lys Val Pro Phe Lys Arg Lys Lys  
 385 390 395 400

Val Val Leu Lys Gln Ile Lys  
 405

<210> 45

<211> 1104

<212> DNA

<213> Streptococcus pneumoniae

<400> 45

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 gagaagattc tatcaaccat tgtttcaaAT ctggatccag aaaagtatga tattgatatt 120  
 cttgaaatgg agcactttga caagggatat gaatctgttc caaagcatgt acgcatttta 180

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aaatcccttc aagattatcg ccaaaccaga tggttacgag cttttttgtg gagaatgaga 240
atattattttc caagactgac tcgtcggttg cttgtaaaag atgattatga tgttgaagtt 300
tctttttacca ttatgaatcc accactgttg ttctctaaaa gaagagaagt caagaagata 360
tcttggattc atggaagtat tgaagaactt ctttaaggata gctctaaaag agaatcacat 420
agaagccagt tggatgctgc gaatacaatt gtagggattt caaaaaagac cagcaattct 480
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tttcagacta ttctagaaaa atctcaagag aagatcgata tcgagattgc tcctcaaagt 600
atctgtacta tcggacggat tgaggaaaat aagggttctg accgtgtagt ggaagtgata 660
cgattattac accaagaggg aaaaaactat catctctatt ttatcggggc tggtgatatg 720
gaagaggaac tgaaaaaacg agtcaaagag tatgggattg aggactatgt acatttcctt 780
ggttatcaaa aaaatcctta tcagtatcta tctcagacga aagttctttt gtctatgtct 840
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tctacggacg ttggaggggc tgaggaatta tcccaagaag gacgatttgg acaaatcatt 960
gagagcaatc aagaggcagc tcaggcgatt actaattaca tgacttctgc ctcaaacttt 1020
gatgtcgatg aggctagcca attcattcaa caatttacia ttacaaaaca aatcgaacaa 1080
gtagaaaaaac tattagagga gtag                                     1104

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<210> 46

<211> 367

<212> PRT

<213> Streptococcus pneumoniae

<400> 46

```

Met Val Ala Lys Lys Lys Ile Leu Phe Phe Met Trp Ser Phe Ser Leu
  1             5             10             15

```

```

Gly Gly Gly Ala Glu Lys Ile Leu Ser Thr Ile Val Ser Asn Leu Asp
      20             25             30

```

```

Pro Glu Lys Tyr Asp Ile Asp Ile Leu Glu Met Glu His Phe Asp Lys
      35             40             45

```

```

Gly Tyr Glu Ser Val Pro Lys His Val Arg Ile Leu Lys Ser Leu Gln
      50             55             60

```

```

Asp Tyr Arg Gln Thr Arg Trp Leu Arg Ala Phe Leu Trp Arg Met Arg
      65             70             75             80

```

Ile Tyr Phe Pro Arg Leu Thr Arg Arg Leu Leu Val Lys Asp Asp Tyr  
                   85                  90                  95

Asp Val Glu Val Ser Phe Thr Ile Met Asn Pro Pro Leu Leu Phe Ser  
                   100                  105                  110

Lys Arg Arg Glu Val Lys Lys Ile Ser Trp Ile His Gly Ser Ile Glu  
                   115                  120                  125

Glu Leu Leu Lys Asp Ser Ser Lys Arg Glu Ser His Arg Ser Gln Leu  
                   130                  135                  140

Asp Ala Ala Asn Thr Ile Val Gly Ile Ser Lys Lys Thr Ser Asn Ser  
                   145                  150                  155                  160

Ile Lys Glu Val Tyr Pro Asp Tyr Thr Ser Lys Leu Gln Thr Ile Tyr  
                   165                  170                  175

Asn Gly Tyr Asp Phe Gln Thr Ile Leu Glu Lys Ser Gln Glu Lys Ile  
                   180                  185                  190

Asp Ile Glu Ile Ala Pro Gln Ser Ile Cys Thr Ile Gly Arg Ile Glu  
                   195                  200                  205

Glu Asn Lys Gly Ser Asp Arg Val Val Glu Val Ile Arg Leu Leu His  
                   210                  215                  220

Gln Glu Gly Lys Asn Tyr His Leu Tyr Phe Ile Gly Ala Gly Asp Met  
                   225                  230                  235                  240

Glu Glu Glu Leu Lys Lys Arg Val Lys Glu Tyr Gly Ile Glu Asp Tyr  
                   245                  250                  255

Val His Phe Leu Gly Tyr Gln Lys Asn Pro Tyr Gln Tyr Leu Ser Gln  
                   260                  265                  270

Thr Lys Val Leu Leu Ser Met Ser Lys Gln Glu Gly Phe Pro Gly Val  
                   275                  280                  285

Tyr Val Glu Ala Leu Ser Leu Gly Leu Pro Phe Ile Ser Thr Asp Val  
 290 295 300

Gly Gly Ala Glu Glu Leu Ser Gln Glu Gly Arg Phe Gly Gln Ile Ile  
 305 310 315 320

Glu Ser Asn Gln Glu Ala Ala Gln Ala Ile Thr Asn Tyr Met Thr Ser  
 325 330 335

Ala Ser Asn Phe Asp Val Asp Glu Ala Ser Gln Phe Ile Gln Gln Phe  
 340 345 350

Thr Ile Thr Lys Gln Ile Glu Gln Val Glu Lys Leu Leu Glu Glu  
 355 360 365

<210> 47

<211> 987

<212> DNA

<213> Streptococcus pneumoniae

<400> 47

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 gatggtgcaa cagatgaaag tggctcgcttg tgtgattcaa tcgctgaaca agatgacagg 180  
 gtgtcagtgcc ttcataaaaa gaacgaagga ttgtcgcaag cacgaaatga tgggatgaag 240  
 caggctcacg gggattatct gattttttatt gactcagatg attatatcca tccagaaatg 300  
 attcagagct tatatgagca attagttcaa gaagatgcgg atgttttcgag ctgtggtgtc 360  
 atgaatgtct atgctaataga tgaaagccca cagtcagcca atcaggatga ctattttgtc 420  
 tgtgattctc aaacatttct aaaggaatac ctcataggtg aaaaaatacc tgggacgatt 480  
 tgcaataagc taatcaagag acagattgca actgccttat ccttttcctaa ggggttgatt 540  
 tacgaagatg cctattacca ttttgattta atcaagttgg ccaagaagta tgtggttaat 600  
 actaaaccct attattacta tttccataga ggggatagta ttacgaccaa accctatgca 660  
 gagaaggatt tagcctatat tgatatctac caaaagtttt ataatagaagt tgtgaaaaac 720  
 tatcctgact tgaaagaggt cgcttttttc agattggcct atgcccactt ctttattctg 780  
 gataagatgt tgctagatga tcagtataaa cagtttgaag cctattctca gattcatcgt 840  
 tttttaaaag gccatgcctt tgctatttct aggaatccaa ttttccgtaa ggggagaaga 900



attagtgcctt tggccctatt cataaatatt tccttatatc gattccttatt actgaaaaat 960  
 attgaaaaat ctaaaaaatt acattag 987

<210> 48

<211> 328

<212> PRT

<213> Streptococcus pneumoniae

<400> 48

Met Glu Thr Ala Leu Ile Ser Val Ile Val Pro Val Tyr Asn Val Ala  
 1 5 10 15

Gln Tyr Leu Glu Lys Ser Ile Ala Ser Ile Gln Lys Gln Thr Tyr Gln  
 20 25 30

Asn Leu Glu Ile Ile Leu Val Asp Asp Gly Ala Thr Asp Glu Ser Gly  
 35 40 45

Arg Leu Cys Asp Ser Ile Ala Glu Gln Asp Asp Arg Val Ser Val Leu  
 50 55 60

His Lys Lys Asn Glu Gly Leu Ser Gln Ala Arg Asn Asp Gly Met Lys  
 65 70 75 80

Gln Ala His Gly Asp Tyr Leu Ile Phe Ile Asp Ser Asp Asp Tyr Ile  
 85 90 95

His Pro Glu Met Ile Gln Ser Leu Tyr Glu Gln Leu Val Gln Glu Asp  
 100 105 110

Ala Asp Val Ser Ser Cys Gly Val Met Asn Val Tyr Ala Asn Asp Glu  
 115 120 125

Ser Pro Gln Ser Ala Asn Gln Asp Asp Tyr Phe Val Cys Asp Ser Gln  
 130 135 140

Thr Phe Leu Lys Glu Tyr Leu Ile Gly Glu Lys Ile Pro Gly Thr Ile  
 145 150 155 160

Cys Asn Lys Leu Ile Lys Arg Gln Ile Ala Thr Ala Leu Ser Phe Pro  
 165 170 175

Lys Gly Leu Ile Tyr Glu Asp Ala Tyr Tyr His Phe Asp Leu Ile Lys  
 180 185 190

Leu Ala Lys Lys Tyr Val Val Asn Thr Lys Pro Tyr Tyr Tyr Tyr Phe  
 195 200 205

His Arg Gly Asp Ser Ile Thr Thr Lys Pro Tyr Ala Glu Lys Asp Leu  
 210 215 220

Ala Tyr Ile Asp Ile Tyr Gln Lys Phe Tyr Asn Glu Val Val Lys Asn  
 225 230 235 240

Tyr Pro Asp Leu Lys Glu Val Ala Phe Phe Arg Leu Ala Tyr Ala His  
 245 250 255

Phe Phe Ile Leu Asp Lys Met Leu Leu Asp Asp Gln Tyr Lys Gln Phe  
 260 265 270

Glu Ala Tyr Ser Gln Ile His Arg Phe Leu Lys Gly His Ala Phe Ala  
 275 280 285

Ile Ser Arg Asn Pro Ile Phe Arg Lys Gly Arg Arg Ile Ser Ala Leu  
 290 295 300

Ala Leu Phe Ile Asn Ile Ser Leu Tyr Arg Phe Leu Leu Leu Lys Asn  
 305 310 315 320

Ile Glu Lys Ser Lys Lys Leu His  
 325

&lt;210&gt; 49

&lt;211&gt; 735

&lt;212&gt; DNA

<213> *Streptococcus pneumoniae*

&lt;400&gt; 49

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tatggtcatt caatcattct caaagatata aattttgcac ttaacaaggg tgaaattggt 120
gggtctagcag ggagaaatgg agttggtaag agtacgttga tgaaaattct tgttcagaat 180
aatcaaccga cttcaggtaa tattataagc agtgataatg ttgggtatit aatcgaagaa 240
ccaaaattat ttttatctaa aacagggtta gagaatttaa aatatttgtc aaatttatat 300
gggtgttgact acaatcaaga aagattttaga tgtttgatcc aagagtttaga tttgactcag 360
tctattaata aaaaagtaaa gacctattct ttgggtacaa aacaaaaatt agctttgctt 420
ctaactctcg ttacggaacc tgatatattg atttttagatg aaccgactaa tggtttagat 480
attgaatcat cacaaatagt ttttagcggtt ctaaaaaaat tagctttaca tgaaaatgtg 540
ggaattttta tatcgagtca taaattagaa gacattgaag aaatttgtga gagagttctt 600
ttcttgagaa acgggctttt gacatttcaa aaagtaggaa aagatagtca taatttcttg 660
tttgagatag ctttttcatc agctacagat agagacattt tcattaccaa acaagaattt 720
tgggatattg tttag                                     735

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&lt;210&gt; 50

&lt;211&gt; 244

&lt;212&gt; PRT

<213> *Streptococcus pneumoniae*

&lt;400&gt; 50

```

Met Arg Ile Lys Glu Lys Thr Asn Asn Ile Asn Gly Gly Ile Lys Asn
  1              5              10              15

Val Ser Lys His Tyr Gly His Ser Ile Ile Leu Lys Asp Ile Asn Phe
          20              25              30

Ala Leu Asn Lys Gly Glu Ile Val Gly Leu Ala Gly Arg Asn Gly Val
          35              40              45

Gly Lys Ser Thr Leu Met Lys Ile Leu Val Gln Asn Asn Gln Pro Thr
          50              55              60

```

Ser Gly Asn Ile Ile Ser Ser Asp Asn Val Gly Tyr Leu Ile Glu Glu  
65 70 75 80

Pro Lys Leu Phe Leu Ser Lys Thr Gly Leu Glu Asn Leu Lys Tyr Leu  
85 90 95

Ser Asn Leu Tyr Gly Val Asp Tyr Asn Gln Glu Arg Phe Arg Cys Leu  
100 105 110

Ile Gln Glu Leu Asp Leu Thr Gln Ser Ile Asn Lys Lys Val Lys Thr  
115 120 125

Tyr Ser Leu Gly Thr Lys Gln Lys Leu Ala Leu Leu Leu Thr Leu Val  
130 135 140

Thr Glu Pro Asp Ile Leu Ile Leu Asp Glu Pro Thr Asn Gly Leu Asp  
145 150 155 160

Ile Glu Ser Ser Gln Ile Val Leu Ala Val Leu Lys Lys Leu Ala Leu  
165 170 175

His Glu Asn Val Gly Ile Leu Ile Ser Ser His Lys Leu Glu Asp Ile  
180 185 190

Glu Glu Ile Cys Glu Arg Val Leu Phe Leu Glu Asn Gly Leu Leu Thr  
195 200 205

Phe Gln Lys Val Gly Lys Asp Ser His Asn Phe Leu Phe Glu Ile Ala  
210 215 220

Phe Ser Ser Ala Thr Asp Arg Asp Ile Phe Ile Thr Lys Gln Glu Phe  
225 230 235 240

Trp Asp Ile Val

&lt;210&gt; 51

&lt;211&gt; 1704

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 51

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atgactgaat tagataaacg tcaccgcagt agcatttatg acagcatggg taaatcacct 60
aaccgtgcta tgcttcgtgc gactggtatg acagataagg actttgaaac atcgattgtg 120
ggagtgattt cgacttgggc ggaaaataca ccatgtaaca ttcacttgca tgatttcggg 180
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gacatcatcg cggactccat cgaggcggct atgagtggtc acaacgtgga tgccttcgtc 360
gctatcggtg gctgtgacaa gaacatgcct ggatctatga ttgctattgc taatatggat 420
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atggacaagt caggtaaaaa ataa 1704

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&lt;210&gt; 52

&lt;211&gt; 567

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 52

Met Thr Glu Leu Asp Lys Arg His Arg Ser Ser Ile Tyr Asp Ser Met  
 1 5 10 15

Val Lys Ser Pro Asn Arg Ala Met Leu Arg Ala Thr Gly Met Thr Asp  
 20 25 30

Lys Asp Phe Glu Thr Ser Ile Val Gly Val Ile Ser Thr Trp Ala Glu  
 35 40 45

Asn Thr Pro Cys Asn Ile His Leu His Asp Phe Gly Lys Leu Ala Lys  
 50 55 60

Glu Gly Val Lys Ser Ala Gly Ala Trp Pro Val Gln Phe Gly Thr Ile  
 65 70 75 80

Thr Val Ala Asp Gly Ile Ala Met Gly Thr Pro Gly Met Arg Phe Ser  
 85 90 95

Leu Thr Ser Arg Asp Ile Ile Ala Asp Ser Ile Glu Ala Ala Met Ser  
 100 105 110

Gly His Asn Val Asp Ala Phe Val Ala Ile Gly Gly Cys Asp Lys Asn  
 115 120 125

Met Pro Gly Ser Met Ile Ala Ile Ala Asn Met Asp Ile Pro Ala Ile  
 130 135 140

Phe Ala Tyr Gly Gly Thr Ile Ala Pro Gly Asn Leu Asp Gly Lys Asp  
 145 150 155 160

Ile Asp Leu Val Ser Val Phe Glu Gly Ile Gly Lys Trp Asn His Gly  
 165 170 175

Asp Met Thr Ala Glu Asp Val Lys Arg Leu Glu Cys Asn Ala Cys Pro  
 180 185 190

Gly Pro Gly Gly Cys Gly Gly Met Tyr Thr Ala Asn Thr Met Ala Thr  
 195 200 205

Ala Ile Glu Val Leu Gly Met Ser Leu Pro Gly Ser Ser Ser His Pro  
 210 215 220

Ala Glu Ser Ala Asp Lys Lys Glu Asp Ile Glu Ala Ala Gly Arg Ala  
 225 230 235 240

Val Val Lys Met Leu Glu Leu Gly Leu Lys Pro Ser Asp Ile Leu Thr  
 245 250 255

Arg Glu Ala Phe Glu Asp Ala Ile Thr Val Thr Met Ala Leu Gly Gly  
 260 265 270

Ser Thr Asn Ala Thr Leu His Leu Leu Ala Ile Ala His Ala Ala Asn  
 275 280 285

Val Asp Leu Ser Leu Glu Asp Phe Asn Thr Ile Gln Glu Arg Val Pro  
 290 295 300

His Leu Ala Asp Leu Lys Pro Ser Gly Gln Tyr Val Phe Gln Asp Leu  
 305 310 315 320

Tyr Glu Val Gly Gly Val Pro Ala Val Met Lys Tyr Leu Leu Ala Asn  
 325 330 335

Gly Phe Leu His Gly Asp Arg Ile Thr Cys Thr Gly Lys Thr Val Ala  
 340 345 350

Glu Asn Leu Ala Asp Phe Ala Asp Leu Thr Pro Gly Gln Lys Val Ile  
 355 360 365

Met Pro Leu Glu Asn Pro Lys Arg Ala Asp Gly Pro Leu Ile Ile Leu  
 370 375 380

Asn Gly Asn Leu Ala Pro Asp Gly Ala Val Ala Lys Val Ser Gly Val  
 385 390 395 400

Lys Val Arg Arg His Val Gly Pro Ala Lys Val Phe Asp Ser Glu Glu  
 405 410 415

Asp Ala Ile Gln Ala Val Leu Thr Asp Glu Ile Val Asp Gly Asp Val  
 420 425 430

Val Val Val Arg Phe Val Gly Pro Lys Gly Gly Pro Gly Met Pro Glu  
 435 440 445

Met Leu Ser Leu Ser Ser Met Ile Val Gly Lys Gly Gln Gly Asp Lys  
 450 455 460

Val Ala Leu Leu Thr Asp Gly Arg Phe Ser Gly Gly Thr Tyr Gly Leu  
 465 470 475 480

Val Val Gly His Ile Ala Pro Glu Ala Gln Asp Gly Gly Pro Ile Ala  
 485 490 495

Tyr Leu Arg Thr Gly Asp Ile Val Thr Val Asp Gln Asp Thr Lys Glu  
 500 505 510

Ile Ser Met Ala Val Ser Glu Glu Glu Leu Glu Lys Arg Lys Ala Glu  
 515 520 525

Thr Thr Leu Pro Pro Leu Tyr Ser Arg Gly Val Leu Gly Lys Tyr Ala  
 530 535 540

His Ile Val Ser Ser Ala Ser Arg Gly Ala Val Thr Asp Phe Trp Asn  
 545 550 555 560

Met Asp Lys Ser Gly Lys Lys  
 565



&lt;210&gt; 53

&lt;211&gt; 274

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 53

```

atgttataat aaaaataaag aatttaagga gaaatacaat atgtcaattt ttattggagg 60
agcatggcca tatgcaaacg gttcgttaca tattggtcac gcggcagcgc ttttaccggg 120
ggatattcctt gcaagatact atcgtcagaa gggagaggaa gttttatatg tttctggaag 180
tgattgtaat ggaaccccta tttctatcag agctaaaaaa gaaaataagt ctgtgaaaga 240
aattgctgat ttttatcata aggaatttaa tcca                                274

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&lt;210&gt; 54

&lt;211&gt; 91

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 54

```

Cys Tyr Asn Lys Asn Lys Glu Phe Lys Glu Lys Tyr Asn Met Ser Ile
  1              5              10              15

Phe Ile Gly Gly Ala Trp Pro Tyr Ala Asn Gly Ser Leu His Ile Gly
      20              25              30

His Ala Ala Ala Leu Leu Pro Gly Asp Ile Leu Ala Arg Tyr Tyr Arg
      35              40              45

Gln Lys Gly Glu Glu Val Leu Tyr Val Ser Gly Ser Asp Cys Asn Gly
      50              55              60

Thr Pro Ile Ser Ile Arg Ala Lys Lys Glu Asn Lys Ser Val Lys Glu
      65              70              75              80

Ile Ala Asp Phe Tyr His Lys Glu Phe Asn Pro
      85              90

```

&lt;210&gt; 55

&lt;211&gt; 1065

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 55

```

atgacaacat tattttcaaa aattaaagaa gtaacagaac ttgctgcagt ctcaggatcat 60
gaagcgccctg tccgtgctta tcttcgtgaa aagttgacac cgcattgtgga tgaagtgggtg 120
acagatgggtt tgggtgggtat ttttggtatc aaacattcag aagctgtgga tgcaccgcgc 180
gtcttggtcg cttctcatat ggacgaagtt ggttttatgg tcagcgaaat caagccagat 240
ggtaccttcc gtgtcgtaga aatcgggtggc tggaacccca tgggtgggttag cagccaacgt 300
ttcaaactct tgactcgtga tggatcatgaa attcctgtga tttcagggttc tgttcctccg 360
catttgactc gtggaaaggg gggaccaacc atgccagcca ttgccgatat cgtttttgat 420
ggtgggttttg cggacaaggc tgaggcagaa agttttggca tccgtcctgg tgataccatt 480
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ctcggcaatg aactctatct gggttctaac gtccaagaag aagttggtct gcgtggcgct 660
catacctcta caaccaagtt tgaccagaa gtcttcctcg cagttgattg ctcaccagca 720
ggtgatgtct acgggtgtca aggcaagatt ggagatggaa ccttgattcg tttctatgat 780
ccaggtcact tgcttctccc agggatgaag gatttccttt tgacaacggc tgaagaagct 840
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caaaccctct atgcaatgga tgacttcta gaagcgcaag ctttcttaca agccttggtg 1020
aagaaattgg atcgttcaac ggttgatttg attaaacatt attaa 1065

```

&lt;210&gt; 56

&lt;211&gt; 354

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 56

```

Met Thr Thr Leu Phe Ser Lys Ile Lys Glu Val Thr Glu Leu Ala Ala
  1              5              10              15

Val Ser Gly His Glu Ala Pro Val Arg Ala Tyr Leu Arg Glu Lys Leu
 20              25              30

```

Thr Pro His Val Asp Glu Val Val Thr Asp Gly Leu Gly Gly Ile Phe  
 35 40 45

Gly Ile Lys His Ser Glu Ala Val Asp Ala Pro Arg Val Leu Val Ala  
 50 55 60

Ser His Met Asp Glu Val Gly Phe Met Val Ser Glu Ile Lys Pro Asp  
 65 70 75 80

Gly Thr Phe Arg Val Val Glu Ile Gly Gly Trp Asn Pro Met Val Val  
 85 90 95

Ser Ser Gln Arg Phe Lys Leu Leu Thr Arg Asp Gly His Glu Ile Pro  
 100 105 110

Val Ile Ser Gly Ser Val Pro Pro His Leu Thr Arg Gly Lys Gly Gly  
 115 120 125

Pro Thr Met Pro Ala Ile Ala Asp Ile Val Phe Asp Gly Gly Phe Ala  
 130 135 140

Asp Lys Ala Glu Ala Glu Ser Phe Gly Ile Arg Pro Gly Asp Thr Ile  
 145 150 155 160

Val Pro Asp Ser Ser Ala Ile Leu Thr Ala Asn Glu Lys Asn Ile Ile  
 165 170 175

Ser Lys Ala Trp Asp Asn Arg Tyr Gly Val Leu Met Val Ser Glu Leu  
 180 185 190

Ala Glu Ala Leu Ser Gly Gln Lys Leu Gly Asn Glu Leu Tyr Leu Gly  
 195 200 205

Ser Asn Val Gln Glu Glu Val Gly Leu Arg Gly Ala His Thr Ser Thr  
 210 215 220

Thr Lys Phe Asp Pro Glu Val Phe Leu Ala Val Asp Cys Ser Pro Ala  
 225 230 235 240

Gly Asp Val Tyr Gly Gly Gln Gly Lys Ile Gly Asp Gly Thr Leu Ile  
 245 250 255

Arg Phe Tyr Asp Pro Gly His Leu Leu Leu Pro Gly Met Lys Asp Phe  
 260 265 270

Leu Leu Thr Thr Ala Glu Glu Ala Gly Ile Lys Tyr Gln Tyr Tyr Cys  
 275 280 285

Gly Lys Gly Gly Thr Asp Ala Gly Ala Ala His Leu Lys Asn Gly Gly  
 290 295 300

Val Pro Ser Thr Thr Ile Gly Val Cys Ala Arg Tyr Ile His Ser His  
 305 310 315 320

Gln Thr Leu Tyr Ala Met Asp Asp Phe Leu Glu Ala Gln Ala Phe Leu  
 325 330 335

Gln Ala Leu Val Lys Lys Leu Asp Arg Ser Thr Val Asp Leu Ile Lys  
 340 345 350

His Tyr

<210> 57

<211> 1182

<212> DNA

<213> Streptococcus pneumoniae

<400> 57

atggaatttt ctatgaaatc agtcaaagga ctactcttta tcatagctag ttttatcttg 60  
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 agcctatctc tgacttttat cctagccact cgtctccac tactagaaag ctggtttcac 180  
 agtttggaga aggtctacac cgtccacaaa ttcacagcct ttctctcaat catcctacta 240  
 atctttcata acttttagtat gggcggtttg tggggctctc gcttagctgc tcagtttggc 300  
 aatcttgcca tctatatctt tgccagcatc atccttgctg cctatttagg caaatacatc 360

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caatacgaag cttggcgatg gattcaccgc ctgggtttacc tagcctatat tttaggactc 420
tttcacatct acatgataat gggcaatcgt ctoccttacat ttaatcttct aagttttctt 480
gttggtagct atgccctttt aggcttacta gctgggtttt atatcatttt tctatatcaa 540
aagatttcct tcccctatct agggaaaatt acccatctca aacgcttaaa tcacgatact 600
agagaaattc aaatccatct tagcagacct ttcaactatc aatcaggaca atttgccttt 660
ctaaagattt tccaagaagg ctttgaaagt gctccgcac ccttttctat ctcaggaggt 720
catggtcaaa ctctttactt tactgttaaa acttcaggcg accataccaa gaatatctat 780
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<210> 58

<211> 394

<212> PRT

<213> Streptococcus pneumoniae

<400> 58

Met Glu Phe Ser Met Lys Ser Val Lys Gly Leu Leu Phe Ile Ile Ala

1

5

10

15

Ser Phe Ile Leu Thr Leu Leu Thr Trp Met Asn Thr Ser Pro Gln Phe

20

25

30

Met Ile Pro Gly Leu Ala Leu Thr Ser Leu Ser Leu Thr Phe Ile Leu

35

40

45

Ala Thr Arg Leu Pro Leu Leu Glu Ser Trp Phe His Ser Leu Glu Lys

50

55

60

Val Tyr Thr Val His Lys Phe Thr Ala Phe Leu Ser Ile Ile Leu Leu

65

70

75

80

81

Ile Phe His Asn Phe Ser Met Gly Gly Leu Trp Gly Ser Arg Leu Ala  
85 90 95

Ala Gln Phe Gly Asn Leu Ala Ile Tyr Ile Phe Ala Ser Ile Ile Leu  
100 105 110

Val Ala Tyr Leu Gly Lys Tyr Ile Gln Tyr Glu Ala Trp Arg Trp Ile  
115 120 125

His Arg Leu Val Tyr Leu Ala Tyr Ile Leu Gly Leu Phe His Ile Tyr  
130 135 140

Met Ile Met Gly Asn Arg Leu Leu Thr Phe Asn Leu Leu Ser Phe Leu  
145 150 155 160

Val Gly Ser Tyr Ala Leu Leu Gly Leu Leu Ala Gly Phe Tyr Ile Ile  
165 170 175

Phe Leu Tyr Gln Lys Ile Ser Phe Pro Tyr Leu Gly Lys Ile Thr His  
180 185 190

Leu Lys Arg Leu Asn His Asp Thr Arg Glu Ile Gln Ile His Leu Ser  
195 200 205

Arg Pro Phe Asn Tyr Gln Ser Gly Gln Phe Ala Phe Leu Lys Ile Phe  
210 215 220

Gln Glu Gly Phe Glu Ser Ala Pro His Pro Phe Ser Ile Ser Gly Gly  
225 230 235 240

His Gly Gln Thr Leu Tyr Phe Thr Val Lys Thr Ser Gly Asp His Thr  
245 250 255

Lys Asn Ile Tyr Asp Asn Leu Gln Ala Gly Ser Lys Val Thr Leu Asp  
260 265 270

Arg Ala Tyr Gly His Met Ile Ile Glu Glu Gly Arg Glu Asn Gln Val  
275 280 285

Trp Ile Ala Gly Gly Ile Gly Ile Thr Pro Phe Ile Ser Tyr Ile Arg  
 290 295 300

Glu His Pro Ile Leu Asp Lys Gln Val His Phe Tyr Tyr Ser Phe Arg  
 305 310 315 320

Gly Asp Glu Asn Ala Val Tyr Leu Asp Leu Leu Arg Asn Tyr Ala Gln  
 325 330 335

Lys Asn Pro Asn Phe Glu Leu His Leu Ile Asp Ser Thr Lys Asp Gly  
 340 345 350

Tyr Leu Asn Phe Glu Gln Lys Glu Val Pro Glu His Ala Thr Val Tyr  
 355 360 365

Met Cys Gly Pro Ile Ser Met Met Lys Ala Leu Ala Lys Gln Ile Lys  
 370 375 380

Lys Gln Asn Pro Lys Thr Glu His Ile Tyr  
 385 390

<210> 59

<211> 900

<212> DNA

<213> Streptococcus pneumoniae

<400> 59

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 cgcaataaaa tcatgggaat ttacacgact gataaggagc aaattgtctt tatcgacaca 180  
 ccagggattc acaagcctaa aacagctctc ggagatttca tggttgagtc tgcctacagt 240  
 acccttcgcg aagtggacac tgtttctttc atggtgcctg ctgatgaagc gcgtggtaag 300  
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 aataaaatcg ataaggtcca tccagaccag ctcttgtctc agattgatga cttccgtaat 420  
 caaatggact ttaaggaaat tgttccaatc tcagcccttc agggaaataa cgtgtctcgt 480  
 ctagtggata ttttgagtga aaatctggat gaaggtttcc aatatttccc gtctgatcaa 540

atcacagacc atccagaacg tttcttggtt tcagaaatgg ttcgcgagaa agtcttgcac 600  
 ctaactcgtg aagagattcc gcattctgta gcagtagttg ttgactctat gaaacgagac 660  
 gaagagacag acaaggttca catccgtgca accatcatgg tcgagcgcgga tagccaaaaa 720  
 gggattatca tcggtaaagg tggcgctatg ctttaagaaaa tcggtagcat ggcccgctgt 780  
 gatatcgaac tcatgctagg agacaaggtc ttcctagaaa cctgggtcaa ggtcaagaaa 840  
 aactggcgcg ataaaaagct agatttggct gactttggct ataataaaag agaataactaa 900

<210> 60

<211> 299

<212> PRT

<213> Streptococcus pneumoniae

<400> 60

Met Thr Phe Lys Ser Gly Phe Val Ala Ile Leu Gly Arg Pro Asn Val  
 1 5 10 15

Gly Lys Ser Thr Phe Leu Asn His Val Met Gly Gln Lys Ile Ala Ile  
 20 25 30

Met Ser Asp Lys Ala Gln Thr Thr Arg Asn Lys Ile Met Gly Ile Tyr  
 35 40 45

Thr Thr Asp Lys Glu Gln Ile Val Phe Ile Asp Thr Pro Gly Ile His  
 50 55 60

Lys Pro Lys Thr Ala Leu Gly Asp Phe Met Val Glu Ser Ala Tyr Ser  
 65 70 75 80

Thr Leu Arg Glu Val Asp Thr Val Leu Phe Met Val Pro Ala Asp Glu  
 85 90 95

Ala Arg Gly Lys Gly Asp Asp Met Ile Ile Glu Arg Leu Lys Ala Ala  
 100 105 110

Lys Val Pro Val Ile Leu Val Val Asn Lys Ile Asp Lys Val His Pro  
 115 120 125



Asp Gln Leu Leu Ser Gln Ile Asp Asp Phe Arg Asn Gln Met Asp Phe  
 130 135 140

Lys Glu Ile Val Pro Ile Ser Ala Leu Gln Gly Asn Asn Val Ser Arg  
 145 150 155 160

Leu Val Asp Ile Leu Ser Glu Asn Leu Asp Glu Gly Phe Gln Tyr Phe  
 165 170 175

Pro Ser Asp Gln Ile Thr Asp His Pro Glu Arg Phe Leu Val Ser Glu  
 180 185 190

Met Val Arg Glu Lys Val Leu His Leu Thr Arg Glu Glu Ile Pro His  
 195 200 205

Ser Val Ala Val Val Val Asp Ser Met Lys Arg Asp Glu Glu Thr Asp  
 210 215 220

Lys Val His Ile Arg Ala Thr Ile Met Val Glu Arg Asp Ser Gln Lys  
 225 230 235 240

Gly Ile Ile Ile Gly Lys Gly Gly Ala Met Leu Lys Lys Ile Gly Ser  
 245 250 255

Met Ala Arg Arg Asp Ile Glu Leu Met Leu Gly Asp Lys Val Phe Leu  
 260 265 270

Glu Thr Trp Val Lys Val Lys Lys Asn Trp Arg Asp Lys Lys Leu Asp  
 275 280 285

Leu Ala Asp Phe Gly Tyr Asn Glu Arg Glu Tyr  
 290 295

&lt;210&gt; 61

&lt;211&gt; 855

&lt;212&gt; DNA

<213> *Streptococcus pneumoniae*

&lt;400&gt; 61

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ctgcttcttg tttttacaga aggaggactt atgcctgaat tacctgaggt tgaaaccggt 60
tgctcgtggct tagaaaaatt gattatagga aagaagattt cgagtataga aattcgctac 120
cccaagatga ttaagacgga tttggaagag tttaaaggga aattgcctag tcagattatc 180
gagtcaatgg gacgtcgtgg aaaatatttg cttttttatc tgacagacaa ggtcttgatt 240
tcccatttgc ggatggaggg caagtatttt tactatccag accaaggacc tgaacgcaag 300
catgcccattg ttttctttca ttttgaagat ggtggcacgc ttgtttatga ggatgttcgc 360
aagtttggaa ccatggaact cttggtgcct gaccttttag acgtctactt tattttctaaa 420
aaattaggtc ctgaaccaag cgaacaagac tttgatttac aggtctttca atctgccctt 480
gccaagtcca aaaagcctat caaatcccat ctccatagacc agaccttggg agctggactt 540
ggcaatatct atgtggatga ggttctctgg cgagctcagg ttcattccagc tagaccttcc 600
cagactttga cagcagaaga agcgactgcc attcatgacc agaccattgc tgttttgggc 660
caggctgttg aaaaagggtg ctccaccatt cggacttata ccaatgcctt tggggaagat 720
ggaagcatgc aggactttca tcaggcttat gataagactg gtcaagaatg tgtacgctgt 780
ggtaccatca ttgagaaaat tcaactaggc ggacgtggaa cccacttttg tccaaactgt 840
caaaggaggg actga

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855

&lt;210&gt; 62

&lt;211&gt; 284

&lt;212&gt; PRT

<213> *Streptococcus pneumoniae*

&lt;400&gt; 62

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Met Leu Leu Val Phe Thr Glu Gly Gly Leu Met Pro Glu Leu Pro Glu
  1             5             10             15

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Val Glu Thr Val Cys Arg Gly Leu Glu Lys Leu Ile Ile Gly Lys Lys
      20             25             30

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Ile Ser Ser Ile Glu Ile Arg Tyr Pro Lys Met Ile Lys Thr Asp Leu
    35             40             45

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Glu Glu Phe Gln Arg Glu Leu Pro Ser Gln Ile Ile Glu Ser Met Gly  
 50 55 60

Arg Arg Gly Lys Tyr Leu Leu Phe Tyr Leu Thr Asp Lys Val Leu Ile  
 65 70 75 80

Ser His Leu Arg Met Glu Gly Lys Tyr Phe Tyr Tyr Pro Asp Gln Gly  
 85 90 95

Pro Glu Arg Lys His Ala His Val Phe Phe His Phe Glu Asp Gly Gly  
 100 105 110

Thr Leu Val Tyr Glu Asp Val Arg Lys Phe Gly Thr Met Glu Leu Leu  
 115 120 125

Val Pro Asp Leu Leu Asp Val Tyr Phe Ile Ser Lys Lys Leu Gly Pro  
 130 135 140

Glu Pro Ser Glu Gln Asp Phe Asp Leu Gln Val Phe Gln Ser Ala Leu  
 145 150 155 160

Ala Lys Ser Lys Lys Pro Ile Lys Ser His Leu Leu Asp Gln Thr Leu  
 165 170 175

Val Ala Gly Leu Gly Asn Ile Tyr Val Asp Glu Val Leu Trp Arg Ala  
 180 185 190

Gln Val His Pro Ala Arg Pro Ser Gln Thr Leu Thr Ala Glu Glu Ala  
 195 200 205

Thr Ala Ile His Asp Gln Thr Ile Ala Val Leu Gly Gln Ala Val Glu  
 210 215 220

Lys Gly Gly Ser Thr Ile Arg Thr Tyr Thr Asn Ala Phe Gly Glu Asp  
 225 230 235 240

Gly Ser Met Gln Asp Phe His Gln Val Tyr Asp Lys Thr Gly Gln Glu  
 245 250 255

Cys Val Arg Cys Gly Thr Ile Ile Glu Lys Ile Gln Leu Gly Gly Arg  
 260 265 270

Gly Thr His Phe Cys Pro Asn Cys Gln Arg Arg Asp  
 275 280

<210> 63

<211> 633

<212> DNA

<213> Streptococcus pneumoniae

<400> 63

ttgtccaaac tgtcaaagga gggactgatg ggaaaaatca tcggaatcac tgggggaatt 60  
 gcctctggta agtcaactgt gacaaatddd ctaagacagc aaggctttca agtagtggat 120  
 gccgacgcag tcgtccacca actacagaaa cctgggtggtc gtctgtttga ggctctagta 180  
 cagcactttg ggcaagaaat cattcttgaa aacggagaaac tcaatcgccc tctcctagct 240  
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 atccgtgagg aactgggtac tttagagaaa cagttgggtc agacagaaga gattttcttc 360  
 atggatattc ccctactttt tgagcaggac tacagcgatt ggtttgctga gacttggttg 420  
 gtctatgtgg accgagatgc ccaagtggaa cgcttaatga aaagggaacca gttgtccaaa 480  
 gatgaagctg agtctcgtct ggcagcccag tggcctttag aaaaaaagaa agatttggcc 540  
 agccagggtc ttgataataa tggcaatcag aaccagcttc ttaatcaagt gcatatcctt 600  
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<210> 64

<211> 210

<212> PRT

<213> Streptococcus pneumoniae

<400> 64

Met Ser Lys Leu Ser Lys Glu Gly Leu Met Gly Lys Ile Ile Gly Ile  
 1 5 10 15

Thr Gly Gly Ile Ala Ser Gly Lys Ser Thr Val Thr Asn Phe Leu Arg  
 20 25 30

Gln Gln Gly Phe Gln Val Val Asp Ala Asp Ala Val Val His Gln Leu  
 35 40 45

Gln Lys Pro Gly Gly Arg Leu Phe Glu Ala Leu Val Gln His Phe Gly  
 50 55 60

Gln Glu Ile Ile Leu Glu Asn Gly Glu Leu Asn Arg Pro Leu Leu Ala  
 65 70 75 80

Ser Leu Ile Phe Ser Asn Pro Asp Glu Arg Glu Trp Ser Lys Gln Ile  
 85 90 95

Gln Gly Glu Ile Ile Arg Glu Glu Leu Ala Thr Leu Arg Glu Gln Leu  
 100 105 110

Ala Gln Thr Glu Glu Ile Phe Phe Met Asp Ile Pro Leu Leu Phe Glu  
 115 120 125

Gln Asp Tyr Ser Asp Trp Phe Ala Glu Thr Trp Leu Val Tyr Val Asp  
 130 135 140

Arg Asp Ala Gln Val Glu Arg Leu Met Lys Arg Asp Gln Leu Ser Lys  
 145 150 155 160

Asp Glu Ala Glu Ser Arg Leu Ala Ala Gln Trp Pro Leu Glu Lys Lys  
 165 170 175

Lys Asp Leu Ala Ser Gln Val Leu Asp Asn Asn Gly Asn Gln Asn Gln  
 180 185 190

Leu Leu Asn Gln Val His Ile Leu Leu Glu Gly Gly Arg Gln Asp Asp  
 195 200 205

Arg Asp  
 210

&lt;210&gt; 65

&lt;211&gt; 1269

&lt;212&gt; DNA

<213> *Streptococcus pneumoniae*

&lt;400&gt; 65

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tttctgacag gagccagtat ttctttgggt gtacctttta tgcccatctt cgtggaaaat 180
ctaggtgtag ggagtcagca agtcgctttt tatgcaggct tagcaatttc tgtctctgct 240
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gaaatctag                                     1269

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&lt;210&gt; 66

&lt;211&gt; 422

&lt;212&gt; PRT

<213> *Streptococcus pneumoniae*

&lt;400&gt; 66

Met Ile Ile Met Ala Ile Arg Thr Ser Phe Leu Ile Lys Cys Ile Ser

1

5

10

15

Phe Leu Arg Glu Val Gly Lys Met Thr Glu Ile Asn Trp Lys Asp Asn  
 20 25 30

Leu Arg Ile Ala Trp Phe Gly Asn Phe Leu Thr Gly Ala Ser Ile Ser  
 35 40 45

Leu Val Val Pro Phe Met Pro Ile Phe Val Glu Asn Leu Gly Val Gly  
 50 55 60

Ser Gln Gln Val Ala Phe Tyr Ala Gly Leu Ala Ile Ser Val Ser Ala  
 65 70 75 80

Ile Ser Ala Ala Leu Phe Ser Pro Ile Trp Gly Ile Leu Ala Asp Lys  
 85 90 95

Tyr Gly Arg Lys Pro Met Met Ile Arg Ala Gly Leu Ala Met Thr Ile  
 100 105 110

Thr Met Gly Gly Leu Ala Phe Val Pro Asn Ile Tyr Trp Leu Ile Phe  
 115 120 125

Leu Arg Leu Leu Asn Gly Val Phe Ala Gly Phe Val Pro Asn Ala Thr  
 130 135 140

Ala Leu Ile Ala Ser Gln Val Pro Lys Glu Lys Ser Gly Ser Ala Leu  
 145 150 155 160

Gly Thr Leu Ser Thr Gly Val Val Ala Gly Thr Leu Thr Gly Pro Phe  
 165 170 175

Ile Gly Gly Phe Ile Ala Glu Leu Phe Gly Ile Arg Thr Val Phe Leu  
 180 185 190

Leu Val Gly Ser Phe Leu Phe Leu Ala Ala Ile Leu Thr Ile Cys Phe  
 195 200 205

Ile Lys Glu Asp Phe Gln Pro Val Ala Lys Glu Lys Ala Ile Pro Thr  
 210 215 220

Lys Glu Leu Phe Thr Ser Val Lys Tyr Pro Tyr Leu Leu Leu Asn Leu  
 225 230 235 240

Phe Leu Thr Ser Phe Val Ile Gln Phe Ser Ala Gln Ser Ile Gly Pro  
 245 250 255

Ile Leu Ala Leu Tyr Val Arg Asp Leu Gly Gln Thr Glu Asn Leu Leu  
 260 265 270

Phe Val Ser Gly Leu Ile Val Ser Ser Met Gly Phe Ser Ser Met Met  
 275 280 285

Ser Ala Gly Val Met Gly Lys Leu Gly Asp Lys Val Gly Asn His Arg  
 290 295 300

Leu Leu Val Val Ala Gln Phe Tyr Ser Val Ile Ile Tyr Leu Leu Cys  
 305 310 315 320

Ala Asn Ala Ser Ser Pro Leu Gln Leu Gly Leu Tyr Arg Phe Leu Phe  
 325 330 335

Gly Leu Gly Thr Gly Ala Leu Ile Pro Gly Val Asn Ala Leu Leu Ser  
 340 345 350

Lys Met Thr Pro Lys Ala Gly Ile Ser Arg Val Phe Ala Phe Asn Gln  
 355 360 365

Val Phe Phe Tyr Leu Gly Gly Val Val Gly Pro Met Ala Gly Ser Ala  
 370 375 380

Val Ala Gly Gln Phe Gly Tyr His Ala Val Phe Tyr Ala Thr Ser Leu  
 385 390 395 400

Cys Val Ala Phe Ser Cys Leu Phe Asn Leu Ile Gln Phe Arg Thr Leu  
 405 410 415

Leu Lys Val Lys Glu Ile  
 420



&lt;210&gt; 67

&lt;211&gt; 1311

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 67

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cgtatTTtatg caacgggtga gtggctcaat cgttctTTta gcatgattga tacaggagga 180
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gaagaagcag atgttatcgt ttttgtcgtg tctggtaagg aaggaattac tgatgcagac 300
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gaaaatactg agaaatactc tgttatgcgt gccatgcgtg ctattgaccg ttcagatgtg 780
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aaagataacc aactatgaa aaactgggaa gaagatatcc gtgagcagtt ccaatacctg 960
ccttacgcac cgattatctt tgtatcagct ttaaccaagc aacgtctcca caaacttcct 1020
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gatgtcatca tggatgccat tgccatcaac ccaacaccga cagacaaagg aaaacgtctc 1140
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gaagaagaac tcatgcactt ttcttacctg cgtttcttgg aaaatcaaT ccgcaaggcc 1260
tttgtTTttg agggaacacc gattcatctc atcgcaagaa aacgcaaata a 1311

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&lt;210&gt; 68

&lt;211&gt; 436

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 68

Met Ala Leu Pro Thr Ile Ala Ile Val Gly Arg Pro Asn Val Gly Lys

1

5

10

15

Ser Thr Leu Phe Asn Arg Ile Ala Gly Glu Arg Ile Ser Ile Val Glu  
 20 25 30

Asp Val Glu Gly Val Thr Arg Asp Arg Ile Tyr Ala Thr Gly Glu Trp  
 35 40 45

Leu Asn Arg Ser Phe Ser Met Ile Asp Thr Gly Gly Ile Asp Asp Val  
 50 55 60

Asp Ala Pro Phe Met Glu Gln Ile Lys His Gln Ala Glu Ile Ala Met  
 65 70 75 80

Glu Glu Ala Asp Val Ile Val Phe Val Val Ser Gly Lys Glu Gly Ile  
 85 90 95

Thr Asp Ala Asp Glu Tyr Val Ala Arg Lys Leu Tyr Lys Thr His Lys  
 100 105 110

Pro Val Ile Leu Ala Val Asn Lys Val Asp Asn Pro Glu Met Arg Asn  
 115 120 125

Asp Ile Tyr Asp Phe Tyr Ala Leu Gly Leu Gly Glu Pro Leu Pro Ile  
 130 135 140

Ser Ser Val His Gly Ile Gly Thr Gly Asp Val Leu Asp Ala Ile Val  
 145 150 155 160

Glu Asn Leu Pro Asn Glu Tyr Glu Glu Glu Asn Pro Asp Val Ile Lys  
 165 170 175

Phe Ser Leu Ile Gly Arg Pro Asn Val Gly Lys Ser Ser Leu Ile Asn  
 180 185 190

Ala Ile Leu Gly Glu Asp Arg Val Ile Ala Ser Pro Val Ala Gly Thr  
 195 200 205

Thr Arg Asp Ala Ile Asp Thr His Phe Thr Asp Thr Asp Gly Gln Glu  
 210 215 220

Phe Thr Met Ile Asp Thr Ala Gly Met Arg Lys Ser Gly Lys Val Tyr  
 225 230 235 240

Glu Asn Thr Glu Lys Tyr Ser Val Met Arg Ala Met Arg Ala Ile Asp  
 245 250 255

Arg Ser Asp Val Val Leu Met Val Ile Asn Ala Glu Glu Gly Ile Arg  
 260 265 270

Glu Tyr Asp Lys Arg Ile Ala Gly Phe Ala His Glu Ala Gly Lys Gly  
 275 280 285

Met Ile Ile Val Val Asn Lys Trp Asp Thr Leu Glu Lys Asp Asn His  
 290 295 300

Thr Met Lys Asn Trp Glu Glu Asp Ile Arg Glu Gln Phe Gln Tyr Leu  
 305 310 315 320

Pro Tyr Ala Pro Ile Ile Phe Val Ser Ala Leu Thr Lys Gln Arg Leu  
 325 330 335

His Lys Leu Pro Glu Met Ile Lys Gln Ile Ser Glu Ser Gln Asn Thr  
 340 345 350

Arg Ile Pro Ser Ala Val Leu Asn Asp Val Ile Met Asp Ala Ile Ala  
 355 360 365

Ile Asn Pro Thr Pro Thr Asp Lys Gly Lys Arg Leu Lys Ile Phe Tyr  
 370 375 380

Ala Thr Gln Val Ala Thr Lys Pro Pro Thr Phe Val Ile Phe Val Asn  
 385 390 395 400

Glu Glu Glu Leu Met His Phe Ser Tyr Leu Arg Phe Leu Glu Asn Gln  
 405 410 415

Ile Arg Lys Ala Phe Val Phe Glu Gly Thr Pro Ile His Leu Ile Ala  
 420 425 430

Arg Lys Arg Lys

435

&lt;210&gt; 69

&lt;211&gt; 714

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 69

```

atgacagaaa ccattaaatt gatgaaggct catacttcag tgcgcagggt taaagagcaa 60
gaaattcccc aagtagactt aaatgagatt ttgacagcag cccagatggc atcatcttgg 120
aagaatttcc aatcctactc tgtgattgtg gtacgaagtc aagagaagaa agatgccttg 180
tatgaattgg tacctcaaga agccattcgc cagtctgctg ttttccttct ctttgtcggg 240
gatttgaacc gagcagaaaa gggagcccg cttcataccg acaccttcca accccaaggt 300
gtggaaggtc tcttgattag ttcggtcgat gcagctcttg ctggacaaaa cgccttggtg 360
gcagctgaaa gcttgggcta tgggtggtgtg attatcggtt tggttcgata caagtctgaa 420
gaagtggcag agctctttaa cctacctgac tacacctatt ctgtctttgg gatggcactg 480
ggtgtgccaa atcaacatca tgatatgaaa ccgagactgc cactagagaa tgttgtcttt 540
gaggaagaat accaagaaca gtcaactgag gcaatccaag cttatgaccg tgttcagggt 600
gactatgctg gggcgctgc gaccacaagc tggagtcagc gcctagcaga acagtttggg 660
caagctgaac caagctcaac tagaaaaaat cttgaacaga agaaattatt gtag      714

```

&lt;210&gt; 70

&lt;211&gt; 237

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 70

Met Thr Glu Thr Ile Lys Leu Met Lys Ala His Thr Ser Val Arg Arg

1

5

10

15

Phe Lys Glu Gln Glu Ile Pro Gln Val Asp Leu Asn Glu Ile Leu Thr

20

25

30

Ala Ala Gln Met Ala Ser Ser Trp Lys Asn Phe Gln Ser Tyr Ser Val  
 35 40 45

Ile Val Val Arg Ser Gln Glu Lys Lys Asp Ala Leu Tyr Glu Leu Val  
 50 55 60

Pro Gln Glu Ala Ile Arg Gln Ser Ala Val Phe Leu Leu Phe Val Gly  
 65 70 75 80

Asp Leu Asn Arg Ala Glu Lys Gly Ala Arg Leu His Thr Asp Thr Phe  
 85 90 95

Gln Pro Gln Gly Val Glu Gly Leu Leu Ile Ser Ser Val Asp Ala Ala  
 100 105 110

Leu Ala Gly Gln Asn Ala Leu Leu Ala Ala Glu Ser Leu Gly Tyr Gly  
 115 120 125

Gly Val Ile Ile Gly Leu Val Arg Tyr Lys Ser Glu Glu Val Ala Glu  
 130 135 140

Leu Phe Asn Leu Pro Asp Tyr Thr Tyr Ser Val Phe Gly Met Ala Leu  
 145 150 155 160

Gly Val Pro Asn Gln His His Asp Met Lys Pro Arg Leu Pro Leu Glu  
 165 170 175

Asn Val Val Phe Glu Glu Glu Tyr Gln Glu Gln Ser Thr Glu Ala Ile  
 180 185 190

Gln Ala Tyr Asp Arg Val Gln Ala Asp Tyr Ala Gly Ala Arg Ala Thr  
 195 200 205

Thr Ser Trp Ser Gln Arg Leu Ala Glu Gln Phe Gly Gln Ala Glu Pro  
 210 215 220

Ser Ser Thr Arg Lys Asn Leu Glu Gln Lys Lys Leu Leu  
 225 230 235

&lt;210&gt; 71

&lt;211&gt; 729

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 71

```

atgacagaaa ttagactaga gcacgtcagt tatgcctatg gtcaggagag gatttttagag 60
gatatcaacc tacaggtgac ttcaggcgaa gtggtttcca tcctaggccc aagtgggtgtt 120
ggaaagacca ccctctttaa tctaatacgt gggatttttag aagttcagtc agggagaatt 180
gtccttgatg gtgaagaaaa tcccaagggg cgcgtgagtt atatgttgca aaaggatctg 240
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acaaccctga tcatcacgca tagtattgag gaggccctca atctcagcga ccgtatctat 600
atcttgaaaa atcgccctgg gcagattggt tcagaaatta aactagattg gtctgaagat 660
gaggacaagg aagtccaaaa gattgcctac aaacgtcaaa ttttggcgga attaggctta 720
gataagtag                                     729

```

&lt;210&gt; 72

&lt;211&gt; 242

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 72

```

Met Thr Glu Ile Arg Leu Glu His Val Ser Tyr Ala Tyr Gly Gln Glu
  1               5               10               15

```

```

Arg Ile Leu Glu Asp Ile Asn Leu Gln Val Thr Ser Gly Glu Val Val
      20               25               30

```

```

Ser Ile Leu Gly Pro Ser Gly Val Gly Lys Thr Thr Leu Phe Asn Leu
      35               40               45

```

Ile Ala Gly Ile Leu Glu Val Gln Ser Gly Arg Ile Val Leu Asp Gly  
 50 55 60

Glu Glu Asn Pro Lys Gly Arg Val Ser Tyr Met Leu Gln Lys Asp Leu  
 65 70 75 80

Leu Leu Glu His Lys Thr Val Leu Gly Asn Ile Ile Leu Pro Leu Leu  
 85 90 95

Ile Gln Lys Val Asp Lys Ala Glu Ala Ile Ser Arg Ala Asp Lys Ile  
 100 105 110

Leu Ala Thr Phe Gln Leu Thr Ala Val Arg Asp Lys Tyr Pro His Glu  
 115 120 125

Leu Ser Gly Gly Met Arg Gln Arg Val Ala Leu Leu Arg Thr Tyr Leu  
 130 135 140

Phe Gly His Lys Leu Phe Leu Leu Asp Glu Ala Phe Ser Ala Leu Asp  
 145 150 155 160

Glu Met Thr Lys Met Glu Leu His Ala Trp Tyr Leu Glu Ile His Lys  
 165 170 175

Gln Leu Gln Leu Thr Thr Leu Ile Ile Thr His Ser Ile Glu Glu Ala  
 180 185 190

Leu Asn Leu Ser Asp Arg Ile Tyr Ile Leu Lys Asn Arg Pro Gly Gln  
 195 200 205

Ile Val Ser Glu Ile Lys Leu Asp Trp Ser Glu Asp Glu Asp Lys Glu  
 210 215 220

Val Gln Lys Ile Ala Tyr Lys Arg Gln Ile Leu Ala Glu Leu Gly Leu  
 225 230 235 240

Asp Lys

&lt;210&gt; 73

&lt;211&gt; 2433

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 73

atgaactatt caaaagcatt gaatgaatgt atcgaaagtg cctacatggg tgctggacat 60  
 tttggagctc gttatctaga gtcgtggcac ttgttgattg ccatgtctaa tcacagttat 120  
 agtgtagcag gggcaacttt aaatgattat ccgtatgaga tggaccgttt agaagaggtg 180  
 gcttttgaac tgactgaaac ggactatagc caggatgaaa cctttacgga attgccgttc 240  
 tcccgtcggt tgcaggttct ttttgatgaa gcagagtatg tagcgtcagt ggtccatgct 300  
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 actcgtatct tggagagggc tgggttttct tatgaagaca agaaagatca ggtcaagatt 420  
 gctgctcttc gtcgaaatct agaagaacgg gcaggctgga ctctgaaga tctcaaggct 480  
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 gtttcaagca ttagccgtgc cattogccgc aaccagtcag ggattcgcag tcataagcgt 1620  
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 gctctggcag aagttctttt tgaogacgaa tcagccctta tccgctttga tatgagttag 1740  
 tatatggaga aatttgcagc tagtcgtctc aacggagctc ctccaggcta tgtaggatat 1800



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gaagaaggtg gggagttgac agagaaggtt cgcaataaac cctattccgt tctcctcttt 1860
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ggtgtcttga cagatagcaa gggacgcaag gtcgattttt caaataccat tatcattatg 1980
acatcgaatc taggtgacac tgcccttcgt gatgataaga ctggttggtt tggggctaag 2040
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aaagcaggcc agttaaatt tgatattgca taa 2433

```

&lt;210&gt; 74

&lt;211&gt; 810

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 74

Met Asn Tyr Ser Lys Ala Leu Asn Glu Cys Ile Glu Ser Ala Tyr Met

1

5

10

15

Val Ala Gly His Phe Gly Ala Arg Tyr Leu Glu Ser Trp His Leu Leu

20

25

30

Ile Ala Met Ser Asn His Ser Tyr Ser Val Ala Gly Ala Thr Leu Asn

35

40

45

Asp Tyr Pro Tyr Glu Met Asp Arg Leu Glu Glu Val Ala Leu Glu Leu

50

55

60

Thr Glu Thr Asp Tyr Ser Gln Asp Glu Thr Phe Thr Glu Leu Pro Phe

65

70

75

80

Ser Arg Arg Leu Gln Val Leu Phe Asp Glu Ala Glu Tyr Val Ala Ser

85

90

95

101

Val Val His Ala Lys Val Leu Gly Thr Glu His Val Leu Tyr Ala Ile  
100 105 110

Leu His Asp Ser Asn Ala Leu Ala Thr Arg Ile Leu Glu Arg Ala Gly  
115 120 125

Phe Ser Tyr Glu Asp Lys Lys Asp Gln Val Lys Ile Ala Ala Leu Arg  
130 135 140

Arg Asn Leu Glu Glu Arg Ala Gly Trp Thr Arg Glu Asp Leu Lys Ala  
145 150 155 160

Leu Arg Gln Arg His Arg Thr Val Ala Asp Lys Gln Asn Ser Met Ala  
165 170 175

Asn Met Met Gly Met Pro Gln Thr Pro Ser Gly Gly Leu Glu Asp Tyr  
180 185 190

Thr His Asp Leu Thr Glu Gln Ala Arg Ser Gly Lys Leu Glu Pro Val  
195 200 205

Ile Gly Arg Asp Lys Glu Ile Ser Arg Met Ile Gln Ile Leu Ser Arg  
210 215 220

Lys Thr Lys Asn Asn Pro Val Leu Val Gly Asp Ala Gly Val Gly Lys  
225 230 235 240

Thr Ala Leu Ala Leu Gly Leu Ala Gln Arg Ile Ala Ser Gly Asp Val  
245 250 255

Pro Ala Glu Met Ala Lys Met Arg Val Leu Glu Leu Asp Leu Met Asn  
260 265 270

Val Val Ala Gly Thr Arg Phe Arg Gly Asp Phe Glu Glu Arg Met Asn  
275 280 285

Asn Ile Ile Lys Asp Ile Glu Glu Asp Gly Gln Val Ile Leu Phe Ile  
290 295 300

Asp Glu Leu His Thr Ile Met Gly Ser Gly Ser Gly Ile Asp Ser Thr  
 305 310 315 320

Leu Asp Ala Ala Asn Ile Leu Lys Pro Ala Leu Ala Arg Gly Thr Leu  
 325 330 335

Arg Thr Val Gly Ala Thr Thr Gln Glu Glu Tyr Gln Lys His Ile Glu  
 340 345 350

Lys Asp Ala Ala Leu Ser Arg Arg Phe Ala Lys Val Thr Ile Glu Glu  
 355 360 365

Pro Ser Val Ala Asp Ser Met Thr Ile Leu Gln Gly Leu Lys Ala Thr  
 370 375 380

Tyr Glu Lys His His Arg Val Gln Ile Thr Asp Glu Ala Val Glu Thr  
 385 390 395 400

Ala Val Lys Met Ala His Arg Tyr Leu Thr Ser Arg His Leu Pro Asp  
 405 410 415

Ser Ala Ile Asp Leu Leu Asp Glu Ala Ala Ala Thr Val Gln Asn Lys  
 420 425 430

Ala Lys His Val Lys Ala Asp Asp Ser Asp Leu Ser Pro Ala Asp Lys  
 435 440 445

Ala Leu Met Asp Gly Lys Trp Lys Gln Ala Ala Gln Leu Ile Ala Lys  
 450 455 460

Glu Glu Glu Val Pro Val Tyr Lys Asp Leu Val Thr Glu Ser Asp Ile  
 465 470 475 480

Leu Thr Thr Leu Ser Arg Leu Ser Gly Ile Pro Val Gln Lys Leu Thr  
 485 490 495

Gln Thr Asp Ala Lys Lys Tyr Leu Asn Leu Glu Ala Glu Leu His Lys  
 500 505 510

Arg Val Ile Gly Gln Asp Gln Ala Val Ser Ser Ile Ser Arg Ala Ile  
 515 520 525

Arg Arg Asn Gln Ser Gly Ile Arg Ser His Lys Arg Pro Ile Gly Ser  
 530 535 540

Phe Met Phe Leu Gly Pro Thr Gly Val Gly Lys Thr Glu Leu Ala Lys  
 545 550 555 560

Ala Leu Ala Glu Val Leu Phe Asp Asp Glu Ser Ala Leu Ile Arg Phe  
 565 570 575

Asp Met Ser Glu Tyr Met Glu Lys Phe Ala Ala Ser Arg Leu Asn Gly  
 580 585 590

Ala Pro Pro Gly Tyr Val Gly Tyr Glu Glu Gly Gly Glu Leu Thr Glu  
 595 600 605

Lys Val Arg Asn Lys Pro Tyr Ser Val Leu Leu Phe Asp Glu Val Glu  
 610 615 620

Lys Ala His Pro Asp Ile Phe Asn Val Leu Leu Gln Val Leu Asp Asp  
 625 630 635 640

Gly Val Leu Thr Asp Ser Lys Gly Arg Lys Val Asp Phe Ser Asn Thr  
 645 650 655

Ile Ile Ile Met Thr Ser Asn Leu Gly Ala Thr Ala Leu Arg Asp Asp  
 660 665 670

Lys Thr Val Gly Phe Gly Ala Lys Asp Ile Arg Phe Asp Gln Glu Asn  
 675 680 685

Met Glu Lys Arg Met Phe Glu Glu Leu Lys Lys Ala Tyr Arg Pro Glu  
 690 695 700

Phe Ile Asn Arg Ile Asp Glu Lys Val Val Phe His Ser Leu Ser Ser  
 705 710 715 720

Asp His Met Gln Glu Val Val Lys Ile Met Val Lys Pro Leu Val Ala  
 725 730 735

Ser Leu Thr Glu Lys Gly Ile Asp Leu Lys Leu Gln Ala Ser Ala Leu  
 740 745 750

Lys Leu Leu Ala Asn Gln Gly Tyr Asp Pro Glu Met Gly Ala Arg Pro  
 755 760 765

Leu Arg Arg Thr Leu Gln Thr Glu Val Glu Asp Lys Leu Ala Glu Leu  
 770 775 780

Leu Leu Lys Gly Asp Leu Val Ala Gly Ser Thr Leu Lys Ile Gly Val  
 785 790 795 800

Lys Ala Gly Gln Leu Lys Phe Asp Ile Ala  
 805 810

<210> 75

<211> 1008

<212> DNA

<213> Streptococcus pneumoniae

<400> 75

atgaagaaaa catggaaagt gtttttaacg cttgtaacag ctcttgtagc tgttgtgctt 60  
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 gactttatcc tagactggac accaaatacc aaccacacag ggctttatgt tgccaaggaa 180  
 aaaggttatt tcaaagaagc tggagtggat gttgatttga aattgccacc agaagaaaagt 240  
 tcttctgact tggttatcaa cggaaaggca ccatttgcag tgtatttcca agactacatg 300  
 gctaagaaat tggaaaaagg agcaggaatc actgccgttg cagctattgt tgaacacaaat 360  
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 aagaaatatg ggacatggaa tgacccaact gaacttgcta tgttgaaaac cttggtagaa 480  
 tctcaagggtg gagactttga gaaggttgaa aaagtaccaa ataacgactc aaactcaatc 540  
 acaccgattg ccaatggcgt ctttgatact gcttggtttt actacggttg ggatgggtatc 600  
 cttgctaaat ctcaagggtg agatgctaac ttcatgtact tgaaagacta tgtcaaggag 660  
 tttgactact attcaccagt tatcatcgca aacaacgact atctgaaaga taacaaagaa 720

gaagctcgca aagtcaccca agccatcaaa aaaggctacc aatatgccat ggaacatcca 780  
gaagaagctg cagatattct catcaagaat gcacctgaac tcaaggaaaa acgtgacttt 840  
gtcatcgaat ctcaaaaata cttgtcaaaa gaatacgcaa gcgacaagga aaaatgggggt 900  
caatttgacg cagctcgctg gaatgctttc tacaaatggg ataaagaaaa tggatcctt 960  
aaagaagact tgacagacaa aggcttcacc aacgaatttg tgaaataa 1008

<210> 76

<211> 335

<212> PRT

<213> Streptococcus pneumoniae

<400> 76

Met Lys Lys Thr Trp Lys Val Phe Leu Thr Leu Val Thr Ala Leu Val  
1 5 10 15

Ala Val Val Leu Val Ala Cys Gly Gln Gly Thr Ala Ser Lys Asp Asn  
20 25 30

Lys Glu Ala Glu Leu Lys Lys Val Asp Phe Ile Leu Asp Trp Thr Pro  
35 40 45

Asn Thr Asn His Thr Gly Leu Tyr Val Ala Lys Glu Lys Gly Tyr Phe  
50 55 60

Lys Glu Ala Gly Val Asp Val Asp Leu Lys Leu Pro Pro Glu Glu Ser  
65 70 75 80

Ser Ser Asp Leu Val Ile Asn Gly Lys Ala Pro Phe Ala Val Tyr Phe  
85 90 95

Gln Asp Tyr Met Ala Lys Lys Leu Glu Lys Gly Ala Gly Ile Thr Ala  
100 105 110

Val Ala Ala Ile Val Glu His Asn Thr Ser Gly Ile Ile Ser Arg Lys  
115 120 125

Ser Asp Asn Val Ser Ser Pro Lys Asp Leu Val Gly Lys Lys Tyr Gly  
 130 135 140

Thr Trp Asn Asp Pro Thr Glu Leu Ala Met Leu Lys Thr Leu Val Glu  
 145 150 155 160

Ser Gln Gly Gly Asp Phe Glu Lys Val Glu Lys Val Pro Asn Asn Asp  
 165 170 175

Ser Asn Ser Ile Thr Pro Ile Ala Asn Gly Val Phe Asp Thr Ala Trp  
 180 185 190

Ile Tyr Tyr Gly Trp Asp Gly Ile Leu Ala Lys Ser Gln Gly Val Asp  
 195 200 205

Ala Asn Phe Met Tyr Leu Lys Asp Tyr Val Lys Glu Phe Asp Tyr Tyr  
 210 215 220

Ser Pro Val Ile Ile Ala Asn Asn Asp Tyr Leu Lys Asp Asn Lys Glu  
 225 230 235 240

Glu Ala Arg Lys Val Ile Gln Ala Ile Lys Lys Gly Tyr Gln Tyr Ala  
 245 250 255

Met Glu His Pro Glu Glu Ala Ala Asp Ile Leu Ile Lys Asn Ala Pro  
 260 265 270

Glu Leu Lys Glu Lys Arg Asp Phe Val Ile Glu Ser Gln Lys Tyr Leu  
 275 280 285

Ser Lys Glu Tyr Ala Ser Asp Lys Glu Lys Trp Gly Gln Phe Asp Ala  
 290 295 300

Ala Arg Trp Asn Ala Phe Tyr Lys Trp Asp Lys Glu Asn Gly Ile Leu  
 305 310 315 320

Lys Glu Asp Leu Thr Asp Lys Gly Phe Thr Asn Glu Phe Val Lys  
 325 330 335

&lt;210&gt; 77

&lt;211&gt; 762

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 77

```

ttgatgagaa acttgagaag tatactgaga cgacacatta gtctattggg ctttctcgga 60
gtattgtcaa tctggcagtt agcagggttt cttaaacttc tccccaaagt tatcctgccg 120
acacctcttg aaattctcca gccctttggt cgtgacagag aatttctctg gcaccatagc 180
tgggcgacct tgagagtggc tttactgggg ctgattttgg gagttttgat tgccctgtctt 240
atggctgtgc tcatggatag tttgacttgg ctcaatgacc tgatttacct tatgatggtg 300
gtcattcaga ccattccgac cattgccata gtccttatcc tggctctgtg gctagggttat 360
gggattttgc ccaagattgt cttgattatc ttaacgacaa ctttcccat catcgtagt 420
attttgacg gttttaggca ttgcgacaag gatatgctga cttgttttag tctgatgcgg 480
gccaagcctt ggcaaatcct gtggcatttt aaaatcccag ttagcctgcc ttacttttat 540
gcaggctctga gggtcagtgt ctctacgcc tttatcacia ctgtggtatc tgagtgggtg 600
ggaggttttg aaggctcttg tgtttatatg attcagtcta aaaaactgtt tcagtatgat 660
accatgtttg ccattattat tctgggtgtc attatcagtc ttttgggtat gaagctggtc 720
gatatcagtg aaaaatatgt gattaaatgg aaacgttcgt ag 762

```

&lt;210&gt; 78

&lt;211&gt; 253

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 78

```

Met Met Arg Asn Leu Arg Ser Ile Leu Arg Arg His Ile Ser Leu Leu
  1               5               10               15

```

```

Gly Phe Leu Gly Val Leu Ser Ile Trp Gln Leu Ala Gly Phe Leu Lys
      20               25               30

```

```

Leu Leu Pro Lys Phe Ile Leu Pro Thr Pro Leu Glu Ile Leu Gln Pro
  35               40               45

```



Phe Val Arg Asp Arg Glu Phe Leu Trp His His Ser Trp Ala Thr Leu  
 50 55 60

Arg Val Ala Leu Leu Gly Leu Ile Leu Gly Val Leu Ile Ala Cys Leu  
 65 70 75 80

Met Ala Val Leu Met Asp Ser Leu Thr Trp Leu Asn Asp Leu Ile Tyr  
 85 90 95

Pro Met Met Val Val Ile Gln Thr Ile Pro Thr Ile Ala Ile Ala Pro  
 100 105 110

Ile Leu Val Leu Trp Leu Gly Tyr Gly Ile Leu Pro Lys Ile Val Leu  
 115 120 125

Ile Ile Leu Thr Thr Thr Phe Pro Ile Ile Val Ser Ile Leu Asp Gly  
 130 135 140

Phe Arg His Cys Asp Lys Asp Met Leu Thr Leu Phe Ser Leu Met Arg  
 145 150 155 160

Ala Lys Pro Trp Gln Ile Leu Trp His Phe Lys Ile Pro Val Ser Leu  
 165 170 175

Pro Tyr Phe Tyr Ala Gly Leu Arg Val Ser Val Ser Tyr Ala Phe Ile  
 180 185 190

Thr Thr Val Val Ser Glu Trp Leu Gly Gly Phe Glu Gly Leu Gly Val  
 195 200 205

Tyr Met Ile Gln Ser Lys Lys Leu Phe Gln Tyr Asp Thr Met Phe Ala  
 210 215 220

Ile Ile Ile Leu Val Ser Ile Ile Ser Leu Leu Gly Met Lys Leu Val  
 225 230 235 240

Asp Ile Ser Glu Lys Tyr Val Ile Lys Trp Lys Arg Ser  
 245 250

&lt;210&gt; 79

&lt;211&gt; 372

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 79

```

ttgattttta atcctatttg ctgtatgata agggaaaaga aaggggacag agatatggct 60
tttaccaata cccacatgcg atctgctagt tttgggtattg ttaccagctt gcctgatgac 120
atcattgact ctttttggtat tatcatcgac catttcttaa aaaatgtctt tgaattggaa 180
gaagaactcg agtttcaatt gcttaataac caaggaaaga ttaccttcca cttttcaagt 240
caacacctcc ctacagccat tgattttgac ttttaaccatc ctttcgaccc tcgttatccc 300
ccaagagtac tgggttttaga catggacggt agagaaacta tcctcctccc agaagaaaat 360
gacctatttt aa 372

```

&lt;210&gt; 80

&lt;211&gt; 123

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 80

```

Met Ile Phe Asn Pro Ile Cys Cys Met Ile Arg Glu Lys Lys Gly Asp
  1             5             10             15

```

```

Arg Asp Met Ala Phe Thr Asn Thr His Met Arg Ser Ala Ser Phe Gly
      20             25             30

```

```

Ile Val Thr Ser Leu Pro Asp Asp Ile Ile Asp Ser Phe Trp Tyr Ile
      35             40             45

```

```

Ile Asp His Phe Leu Lys Asn Val Phe Glu Leu Glu Glu Leu Glu
      50             55             60

```

```

Phe Gln Leu Leu Asn Asn Gln Gly Lys Ile Thr Phe His Phe Ser Ser
      65             70             75             80

```

110

Gln His Leu Pro Thr Ala Ile Asp Phe Asp Phe Asn His Pro Phe Asp  
85 90 95

Pro Arg Tyr Pro Pro Arg Val Leu Val Leu Asp Met Asp Gly Arg Glu  
100 105 110

Thr Ile Leu Leu Pro Glu Glu Asn Asp Leu Phe  
115 120

<210> 81

<211> 1645

<212> DNA

<213> Streptococcus pneumoniae

<400> 81

acagcgggtgt cattotatct attttaagaa aagtaataat caattgttaa aaatagtaaa 60  
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aaaggtgggg acttgctcgg tactattggc aatttcaatt ttgggaagcc aaggtatttt 180  
atcggatgaa gttgttaacta gttottcacc gatggctaca aaagagtctt ctaatgcaat 240  
tactaatgat ttagataatt caccaactgt taatcagaat cgttctgctg aaatgattgc 300  
ctctaattca accactaatg gtttagataa ttcgttaagt gttaatagca tcagctctaa 360  
tggtactatt cgttccaatt cacaattaga caacagaaca gttgaatcta cagtaacatc 420  
tactaatgaa aataagagtt ataaggaaga tggtataagt gacagaatta tcaaaaaaga 480  
atttgaagat actgctttta gtgtaaaaga ttatgggtgca gtaggtgatg ggattcatga 540  
tgatcgacaa gcaattcaag atgcaataga tgctgcagct caagggctag gtggaggaaa 600  
tgtatatattt cctgaaggaa cttattttagt aaaagaaatt gtttttttaa aaagtcatac 660  
acacttagaa ttgaatgaga aagctacaat totaaatggg ataaatatta agaatacccc 720  
ttccattgtt tttatgacag gtttattttac ggatgatggg gcgcaagtag aatggggccc 780  
aacagaagat attagttatt ctggtggtac gattgatatg aacggtgctt tgaatgaaga 840  
aggaactaaa gcaaaaaatc taccacttat aaattcttca ggtgcatttg ctattgggaa 900  
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tcaaattgca ggttcgaaaa atgtattagt tgataattct cgtttttctt ggcaagcctt 1020  
acccaaaacg atgaaggatg ggcaaatcat aagtaaggag agcattcaga ttgaaccatt 1080  
aactagaaaa ggtttttcctt atgccttgaa tgatgatggg aaaaaatctg aaaatgtgac 1140  
tattcaaaat tcctatttttg gcaaaagtga taaatctggg gaattagtaa cagcaattgg 1200  
cacacactat caaacattgt cgacacagaa cccctctaat attaaaattc aaaataatca 1260

ttttgataac atgatgtatg caggtgtacg ttttacagga ttcaactgatg tattaatcaa 1320  
 aggaaatcgc tttgataaga aagttaaagg agagagtgtg cattatcgag aaagcggagc 1380  
 agcttttagta aatgcttata gctataaaaa cactaaagac ctattagatt taaataaaca 1440  
 ggtgggttatc gccgaaaata tattttaatat tgccgatcct aaaacaaaag cgatacgagt 1500  
 tgcaaaagat agtgcagaat gtttaggaaa agtatcagat attactgtaa caaaaaatgt 1560  
 aattaataat aattctaagg aaacagaaca accaaatatt gaattattac gagttagtga 1620  
 taatttagta gtctcagaga atagt 1645

<210> 82

<211> 548

<212> PRT

<213> Streptococcus pneumoniae

<400> 82

Gln Arg Cys His Ser Ile Tyr Phe Lys Lys Ser Asn Asn Gln Leu Leu  
 1 5 10 15

Lys Ile Val Lys Lys Leu Glu Val Leu Met Lys Tyr Phe Val Pro Asn  
 20 25 30

Glu Val Phe Ser Ile Arg Lys Leu Lys Val Gly Thr Cys Ser Val Leu  
 35 40 45

Leu Ala Ile Ser Ile Leu Gly Ser Gln Gly Ile Leu Ser Asp Glu Val  
 50 55 60

Val Thr Ser Ser Ser Pro Met Ala Thr Lys Glu Ser Ser Asn Ala Ile  
 65 70 75 80

Thr Asn Asp Leu Asp Asn Ser Pro Thr Val Asn Gln Asn Arg Ser Ala  
 85 90 95

Glu Met Ile Ala Ser Asn Ser Thr Thr Asn Gly Leu Asp Asn Ser Leu  
 100 105 110

Ser Val Asn Ser Ile Ser Ser Asn Gly Thr Ile Arg Ser Asn Ser Gln  
 115 120 125

Leu Asp Asn Arg Thr Val Glu Ser Thr Val Thr Ser Thr Asn Glu Asn  
 130 135 140

Lys Ser Tyr Lys Glu Asp Val Ile Ser Asp Arg Ile Ile Lys Lys Glu  
 145 150 155 160

Phe Glu Asp Thr Ala Leu Ser Val Lys Asp Tyr Gly Ala Val Gly Asp  
 165 170 175

Gly Ile His Asp Asp Arg Gln Ala Ile Gln Asp Ala Ile Asp Ala Ala  
 180 185 190

Ala Gln Gly Leu Gly Gly Gly Asn Val Tyr Phe Pro Glu Gly Thr Tyr  
 195 200 205

Leu Val Lys Glu Ile Val Phe Leu Lys Ser His Thr His Leu Glu Leu  
 210 215 220

Asn Glu Lys Ala Thr Ile Leu Asn Gly Ile Asn Ile Lys Asn His Pro  
 225 230 235 240

Ser Ile Val Phe Met Thr Gly Leu Phe Thr Asp Asp Gly Ala Gln Val  
 245 250 255

Glu Trp Gly Pro Thr Glu Asp Ile Ser Tyr Ser Gly Gly Thr Ile Asp  
 260 265 270

Met Asn Gly Ala Leu Asn Glu Glu Gly Thr Lys Ala Lys Asn Leu Pro  
 275 280 285

Leu Ile Asn Ser Ser Gly Ala Phe Ala Ile Gly Asn Ser Asn Asn Val  
 290 295 300

Thr Ile Lys Asn Val Thr Phe Lys Asp Ser Tyr Gln Gly His Ala Ile  
 305 310 315 320

Gln Ile Ala Gly Ser Lys Asn Val Leu Val Asp Asn Ser Arg Phe Leu  
 325 330 335

Gly Gln Ala Leu Pro Lys Thr Met Lys Asp Gly Gln Ile Ile Ser Lys  
 340 345 350

Glu Ser Ile Gln Ile Glu Pro Leu Thr Arg Lys Gly Phe Pro Tyr Ala  
 355 360 365

Leu Asn Asp Asp Gly Lys Lys Ser Glu Asn Val Thr Ile Gln Asn Ser  
 370 375 380

Tyr Phe Gly Lys Ser Asp Lys Ser Gly Glu Leu Val Thr Ala Ile Gly  
 385 390 395 400

Thr His Tyr Gln Thr Leu Ser Thr Gln Asn Pro Ser Asn Ile Lys Ile  
 405 410 415

Gln Asn Asn His Phe Asp Asn Met Met Tyr Ala Gly Val Arg Phe Thr  
 420 425 430

Gly Phe Thr Asp Val Leu Ile Lys Gly Asn Arg Phe Asp Lys Lys Val  
 435 440 445

Lys Gly Glu Ser Val His Tyr Arg Glu Ser Gly Ala Ala Leu Val Asn  
 450 455 460

Ala Tyr Ser Tyr Lys Asn Thr Lys Asp Leu Leu Asp Leu Asn Lys Gln  
 465 470 475 480

Val Val Ile Ala Glu Asn Ile Phe Asn Ile Ala Asp Pro Lys Thr Lys  
 485 490 495

Ala Ile Arg Val Ala Lys Asp Ser Ala Glu Cys Leu Gly Lys Val Ser  
 500 505 510

Asp Ile Thr Val Thr Lys Asn Val Ile Asn Asn Asn Ser Lys Glu Thr  
 515 520 525

Glu Gln Pro Asn Ile Glu Leu Leu Arg Val Ser Asp Asn Leu Val Val  
 530 535 540

Ser Glu Asn Ser

545

<210> 83

<211> 324

<212> DNA

<213> Streptococcus pneumoniae

<400> 83

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gtgatgaaag aaactcagct attaaaaggt gttcttgaag gttgtgtctt ggatatgatt 60
gggtcaaaaag agcgggtatgg ttatgagttg gttcagactt tgcgagaggc tggatttgat 120
actatcgttc caggaactat ttatcctttg ttgcaaaagt tagaaaaaaa tcaatggata 180
agaggcgaca tgcgcccgtc gccagatggg ccagatcgga agtatttttc attaatgaaa 240
gaaggagaag agcgtgtctc agtcttttgg caacaatggg acgatttgag tcaaaaagta 300
gaagggatta agaatggggg ttaa                                     324

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<210> 84

<211> 107

<212> PRT

<213> Streptococcus pneumoniae

<400> 84

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Met Met Lys Glu Thr Gln Leu Leu Lys Gly Val Leu Glu Gly Cys Val
  1              5              10              15

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Leu Asp Met Ile Gly Gln Lys Glu Arg Tyr Gly Tyr Glu Leu Val Gln
      20              25              30

```

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Thr Leu Arg Glu Ala Gly Phe Asp Thr Ile Val Pro Gly Thr Ile Tyr
      35              40              45

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Pro Leu Leu Gln Lys Leu Glu Lys Asn Gln Trp Ile Arg Gly Asp Met
      50              55              60

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Arg Pro Ser Pro Asp Gly Pro Asp Arg Lys Tyr Phe Ser Leu Met Lys  
 65 70 75 80

Glu Gly Glu Glu Arg Val Ser Val Phe Trp Gln Gln Trp Asp Asp Leu  
 85 90 95

Ser Gln Lys Val Glu Gly Ile Lys Asn Gly Gly  
 100 105

<210> 85

<211> 816

<212> DNA

<213> Streptococcus pneumoniae

<400> 85

atgaagaaaa tgaagtatta cgaagaaaca agcgctttgc tacatgagtt ttctgaggag 60  
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 gaagactatc tcagagagca gatctatttg atgatgctag atttctcaga agcagaacga 180  
 gatggcatga gtgcagagga ttatctaggt aagaatccta aaaaaataat gaaagagatt 240  
 ctcaagggag cacctcgag ttctatcaaa gagtcccttt tgacgccaat tcttgtcctg 300  
 gcggtattac gttattatca actactaagt gatttttcta aaggtcctct cttaacagtc 360  
 aatttgctca catttttagg gcaacttctt atttttctga ttggatttgg acttgtggcc 420  
 acaattttac gaagaagttt agtccaagat tctcctaaaa tgaaaattgg cacttacatt 480  
 gttgttggga ctatagttct tctagttggt ttaggatatg taggaatggc aagcttcata 540  
 caagaaggag ccttttatat tccggctccc tgggatagtt tgtctgtctt tacgatttcg 600  
 ctagtatatc gtatttggaa ttggaaagaa gcggtctttc gtccatttgt cagtatgatt 660  
 attgcccac ttgtgggtggg ttctctgctc cgttattatg agtggatggg aatttcaaat 720  
 gttttcctta caaaagttat tcctttagct gtcctcttta ttggaatctt tgtcttgttc 780  
 cgtgggttta agaagataaa atggagtga gatatag 816

<210> 86

<211> 271

<212> PRT

<213> Streptococcus pneumoniae



&lt;400&gt; 86

Met Lys Lys Met Lys Tyr Tyr Glu Glu Thr Ser Ala Leu Leu His Glu  
 1 5 10 15

Phe Ser Glu Glu Asn Gln Lys Tyr Phe Glu Glu Leu Trp Glu Ser Phe  
 20 25 30

Asn Leu Ala Gly Phe Leu Tyr Asp Glu Asp Tyr Leu Arg Glu Gln Ile  
 35 40 45

Tyr Leu Met Met Leu Asp Phe Ser Glu Ala Glu Arg Asp Gly Met Ser  
 50 55 60

Ala Glu Asp Tyr Leu Gly Lys Asn Pro Lys Lys Ile Met Lys Glu Ile  
 65 70 75 80

Leu Lys Gly Ala Pro Arg Ser Ser Ile Lys Glu Ser Leu Leu Thr Pro  
 85 90 95

Ile Leu Val Leu Ala Val Leu Arg Tyr Tyr Gln Leu Leu Ser Asp Phe  
 100 105 110

Ser Lys Gly Pro Leu Leu Thr Val Asn Leu Leu Thr Phe Leu Gly Gln  
 115 120 125

Leu Leu Ile Phe Leu Ile Gly Phe Gly Leu Val Ala Thr Ile Leu Arg  
 130 135 140

Arg Ser Leu Val Gln Asp Ser Pro Lys Met Lys Ile Gly Thr Tyr Ile  
 145 150 155 160

Val Val Gly Thr Ile Val Leu Leu Val Val Leu Gly Tyr Val Gly Met  
 165 170 175

Ala Ser Phe Ile Gln Glu Gly Ala Phe Tyr Ile Pro Ala Pro Trp Asp  
 180 185 190

117

Ser Leu Ser Val Phe Thr Ile Ser Leu Val Ile Gly Ile Trp Asn Trp  
195 200 205

Lys Glu Ala Val Phe Arg Pro Phe Val Ser Met Ile Ile Ala His Leu  
210 215 220

Val Val Gly Ser Leu Leu Arg Tyr Tyr Glu Trp Met Gly Ile Ser Asn  
225 230 235 240

Val Phe Leu Thr Lys Val Ile Pro Leu Ala Val Leu Phe Ile Gly Ile  
245 250 255

Phe Val Leu Phe Arg Gly Phe Lys Lys Ile Lys Trp Ser Glu Val  
260 265 270

<210> 87

<211> 348

<212> DNA

<213> Streptococcus pneumoniae

<400> 87

ctgttttttt atttatactc aatgaaaatc aaagagcaaa ctaggaagct agccgcaggt 60  
tgctcaaaac actgttttga ggttgtagac gaaactgacg aagtcagctc aaaacatggt 120  
tttgagggtg tagatgaaac tgacgaagtc agctcaaaac actgttttga ggttgtagat 180  
gaaactgacg aagtcagctc aaaacactgt tttgagggtg tagatgaaac tgacgaagtc 240  
agctcaaaac atgtttttga ggttgtagat gaaactgacg aagtcagtaa ccatacatc 300  
ggtagggcga cgctgacgtg gtttgaagag attttcgaag agtattaa 348

<210> 88

<211> 115

<212> PRT

<213> Streptococcus pneumoniae

<400> 88

Met Phe Phe Tyr Leu Tyr Ser Met Lys Ile Lys Glu Gln Thr Arg Lys  
1 5 10 15

Leu Ala Ala Gly Cys Ser Lys His Cys Phe Glu Val Val Asp Glu Thr  
 20 25 30

Asp Glu Val Ser Ser Lys His Val Phe Glu Val Val Asp Glu Thr Asp  
 35 40 45

Glu Val Ser Ser Lys His Cys Phe Glu Val Val Asp Glu Thr Asp Glu  
 50 55 60

Val Ser Ser Lys His Cys Phe Glu Val Val Asp Glu Thr Asp Glu Val  
 65 70 75 80

Ser Ser Lys His Val Phe Glu Val Val Asp Glu Thr Asp Glu Val Ser  
 85 90 95

Asn His Thr Tyr Gly Arg Ala Thr Leu Thr Trp Phe Glu Glu Ile Phe  
 100 105 110

Glu Glu Tyr  
 115

<210> 89

<211> 1260

<212> DNA

<213> Streptococcus pneumoniae

<400> 89

atgcagaatc tgaaatttgc cttttcatct atcatggctc acaagatgcg ttctttgctt 60  
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 tccctatctc gtcaagtcaa taaagatatg actaaatctc agaaaaatat tagcgtcttt 180  
 ttctctccta aaaaaagtaa agacgggtct tttactcaga aacaatcagc ttttacgggt 240  
 tctggaaagg aagaggaagt tctgttgaa ccgcaaaaac cgcaagaatc ctgggtccaa 300  
 gaggcagcta aactgaaggg agtggatagt tactatgtaa ccaattcaac gaatgccatc 360  
 ttgacctatc aagataaaaa ggttgagaat gctaatttga caggtggaaa cagaacttac 420  
 atggacgctg ttaagaatga aattattgca ggctgtagtc tgagagagca agatttcaaa 480  
 gagtttgcaa gtgtcatttt gctagatgag gaattgtcca ttagtttatt tgaatctcct 540

caagaggcta ttaacaaggt tgtagaagtc aatggattta gttaccgggt cattgggggt 600  
 tatactagtc cggaggctaa aagatcaaaa atatatgggt ttggtggctt gcctattact 660  
 accaatatct cccttgctgc gaattttaat gtagatgaaa tagctaatat tgtctttcga 720  
 gtgaatgata ccagtttaac cccaactctg ggtccagaac tggcacgaaa aatgacagag 780  
 cttgcaggct tacaacaggg agaataccag gtggcagatg agtccgttgt atttgcagaa 840  
 attcaacaat cgttttagttt tatgacgacg attattagtt ccatcgcagg gatttctctc 900  
 tttgttggag gaactggtgt catgaacatc atgctggttt cggtgacaga gcgcactcgt 960  
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 tcaatcccag tcgccctatt tagtcttgca gtttcggcta gtgttggtat gatttttgga 1200  
 gtcttgccag ccaacaaggc atcgaaactt gatccaattg aagcccttcg ttatgaatga 1260

<210> 90

<211> 419

<212> PRT

<213> Streptococcus pneumoniae

<400> 90

Met Gln Asn Leu Lys Phe Ala Phe Ser Ser Ile Met Ala His Lys Met

1

5

10

15

Arg Ser Leu Leu Thr Met Ile Gly Ile Ile Ile Gly Val Ser Ser Val

20

25

30

Val Val Ile Met Ala Leu Gly Asp Ser Leu Ser Arg Gln Val Asn Lys

35

40

45

Asp Met Thr Lys Ser Gln Lys Asn Ile Ser Val Phe Phe Ser Pro Lys

50

55

60

Lys Ser Lys Asp Gly Ser Phe Thr Gln Lys Gln Ser Ala Phe Thr Val

65

70

75

80

Ser Gly Lys Glu Glu Glu Val Pro Val Glu Pro Pro Lys Pro Gln Glu

85

90

95

120

Ser Trp Val Gln Glu Ala Ala Lys Leu Lys Gly Val Asp Ser Tyr Tyr  
100 105 110

Val Thr Asn Ser Thr Asn Ala Ile Leu Thr Tyr Gln Asp Lys Lys Val  
115 120 125

Glu Asn Ala Asn Leu Thr Gly Gly Asn Arg Thr Tyr Met Asp Ala Val  
130 135 140

Lys Asn Glu Ile Ile Ala Gly Arg Ser Leu Arg Glu Gln Asp Phe Lys  
145 150 155 160

Glu Phe Ala Ser Val Ile Leu Leu Asp Glu Glu Leu Ser Ile Ser Leu  
165 170 175

Phe Glu Ser Pro Gln Glu Ala Ile Asn Lys Val Val Glu Val Asn Gly  
180 185 190

Phe Ser Tyr Arg Val Ile Gly Val Tyr Thr Ser Pro Glu Ala Lys Arg  
195 200 205

Ser Lys Ile Tyr Gly Phe Gly Gly Leu Pro Ile Thr Thr Asn Ile Ser  
210 215 220

Leu Ala Ala Asn Phe Asn Val Asp Glu Ile Ala Asn Ile Val Phe Arg  
225 230 235 240

Val Asn Asp Thr Ser Leu Thr Pro Thr Leu Gly Pro Glu Leu Ala Arg  
245 250 255

Lys Met Thr Glu Leu Ala Gly Leu Gln Gln Gly Glu Tyr Gln Val Ala  
260 265 270

Asp Glu Ser Val Val Phe Ala Glu Ile Gln Gln Ser Phe Ser Phe Met  
275 280 285

Thr Thr Ile Ile Ser Ser Ile Ala Gly Ile Ser Leu Phe Val Gly Gly  
290 295 300

121

Thr Gly Val Met Asn Ile Met Leu Val Ser Val Thr Glu Arg Thr Arg  
305 310 315 320

Glu Ile Gly Leu Arg Lys Ala Leu Gly Ala Thr Arg Ala Asn Ile Leu  
325 330 335

Ile Gln Phe Leu Ile Glu Ser Met Ile Leu Thr Leu Leu Gly Gly Leu  
340 345 350

Ile Gly Leu Thr Ile Ala Ser Gly Leu Thr Ala Leu Ala Gly Leu Leu  
355 360 365

Leu Gln Gly Leu Ile Glu Gly Ile Glu Val Gly Val Ser Ile Pro Val  
370 375 380

Ala Leu Phe Ser Leu Ala Val Ser Ala Ser Val Gly Met Ile Phe Gly  
385 390 395 400

Val Leu Pro Ala Asn Lys Ala Ser Lys Leu Asp Pro Ile Glu Ala Leu  
405 410 415

Arg Tyr Glu

<210> 91

<211> 705

<212> DNA

<213> Streptococcus pneumoniae

<400> 91

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atgggaccat ctgggtcttg taagtccact ctgatgaata cgattggcat gttggataca 180  
ccaaccagtg gagaatatta tcttgaaggt caagaagtgg ctgggcttgg tgaaaaacaa 240  
ctagctaagg tccgtaacca acaaatcggt tttgtctttc agcagttctt tcttctatcg 300  
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aaacgtcgca agttggctga ggaatattta gacaagggtg aattgacaga acgtagtcac 420  
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 ggtaacaaaa ttatgcaatt attggttgat ttgaataaag aaggaaaaac cattatcatg 600  
 gtaacgcatg agcctgagat tgctgcctat gccaaacgtc agattgtcat tcgggatggg 660  
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<210> 92

<211> 234

<212> PRT

<213> Streptococcus pneumoniae

<400> 92

Met Met Lys Gln Leu Ile Ser Leu Lys Asn Ile Phe Arg Ser Tyr Arg  
 1 5 10 15

Asn Gly Asp Gln Glu Leu Gln Val Leu Lys Asn Ile Asn Leu Glu Val  
 20 25 30

Asn Glu Gly Glu Phe Val Ala Ile Met Gly Pro Ser Gly Ser Gly Lys  
 35 40 45

Ser Thr Leu Met Asn Thr Ile Gly Met Leu Asp Thr Pro Thr Ser Gly  
 50 55 60

Glu Tyr Tyr Leu Glu Gly Gln Glu Val Ala Gly Leu Gly Glu Lys Gln  
 65 70 75 80

Leu Ala Lys Val Arg Asn Gln Gln Ile Gly Phe Val Phe Gln Gln Phe  
 85 90 95

Phe Leu Leu Ser Lys Leu Asn Ala Leu Gln Asn Val Glu Leu Pro Leu  
 100 105 110

Ile Tyr Ala Gly Val Ser Ser Ser Lys Arg Arg Lys Leu Ala Glu Glu  
 115 120 125

123

Tyr Leu Asp Lys Val Glu Leu Thr Glu Arg Ser His His Leu Pro Ser  
130 135 140

Glu Leu Ser Gly Gly Gln Lys Gln Arg Val Ala Ile Ala Arg Ala Leu  
145 150 155 160

Val Asn Asn Pro Ser Ile Ile Leu Ala Asp Glu Pro Thr Gly Ala Leu  
165 170 175

Asp Thr Lys Thr Gly Asn Gln Ile Met Gln Leu Leu Val Asp Leu Asn  
180 185 190

Lys Glu Gly Lys Thr Ile Ile Met Val Thr His Glu Pro Glu Ile Ala  
195 200 205

Ala Tyr Ala Lys Arg Gln Ile Val Ile Arg Asp Gly Val Ile Ser Ser  
210 215 220

Asp Ser Ala Gln Leu Gly Lys Glu Glu Asn  
225 230

<210> 93

<211> 1200

<212> DNA

<213> Streptococcus pneumoniae

<400> 93

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ctaaaagatg agcctactca tcttggtgtt gccaaaggaag gaagcgtggc ctctctgtt 180  
ttattgtcag ggacagtaac agcaaaaaat gaacaatatg tttattttga tgctagtaag 240  
ggtgatttag atgaaatcct tgtttctgtg ggcgataagg tcagcgaagg gcaggcttta 300  
gtcaagtaca gtagttcaga agcgcaggcg gcctatgatt cagctagtcg agcagtagct 360  
agggcagatc gtcatatcaa tgaactcaat caagcacgaa atgaagccgc ttcagctccg 420  
gctccacagt taccagcgcc agtaggagga gaagatgcaa cggtgcaaag cccaactcca 480  
gtggctggaa attctgttgc ttctattgac gctcaattgg gtgatgcccg tgatgcgcgt 540



gcagatgctg cggcgcaatt aagcaaggct caaagtcaat tggatgcaac aactgttctc 600  
 agtaccctag agggaaactgt ggtcgaagtc aatagcaatg tttctaaatc tccaacaggg 660  
 gcgagtcaag ttatggttca tattgtcagc aatgaaaatt tacaagtcaa gggagaattg 720  
 tctgagtaca atctagccaa cctttctgta ggtcaagaag taagctttac ttctaaagtg 780  
 tatectgata aaaaatggac tgggaaatta agctatatct ctgactatcc taaaaacaat 840  
 ggtgaagcag ctagtccagc agccgggaat aatacagggt ctaaataccc ttatactatt 900  
 gatgtgacag gcgaggttgg tgatttgaaa caagggtttt ctgtcaacat tgagggttaa 960  
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 gctgacgcag aaaatcaaga aatcacttct ggtttaacga acggtgctaa ggtcatcagt 1140  
 aatccaacat cttccttgga agaaggaaaa gaggtgaagg ctgatgaagc aactaattag 1200

<210> 94

<211> 399

<212> PRT

<213> Streptococcus pneumoniae

<400> 94

Met Lys Lys Lys Asn Gly Lys Ala Lys Lys Trp Gln Leu Tyr Ala Ala  
 1 5 10 15

Ile Gly Ala Ala Ser Val Val Val Leu Gly Ala Gly Gly Ile Leu Leu  
 20 25 30

Phe Arg Gln Pro Ser Gln Thr Ala Leu Lys Asp Glu Pro Thr His Leu  
 35 40 45

Val Val Ala Lys Glu Gly Ser Val Ala Ser Ser Val Leu Leu Ser Gly  
 50 55 60

Thr Val Thr Ala Lys Asn Glu Gln Tyr Val Tyr Phe Asp Ala Ser Lys  
 65 70 75 80

Gly Asp Leu Asp Glu Ile Leu Val Ser Val Gly Asp Lys Val Ser Glu  
 85 90 95

125

Gly Gln Ala Leu Val Lys Tyr Ser Ser Ser Glu Ala Gln Ala Ala Tyr  
100 105 110

Asp Ser Ala Ser Arg Ala Val Ala Arg Ala Asp Arg His Ile Asn Glu  
115 120 125

Leu Asn Gln Ala Arg Asn Glu Ala Ala Ser Ala Pro Ala Pro Gln Leu  
130 135 140

Pro Ala Pro Val Gly Gly Glu Asp Ala Thr Val Gln Ser Pro Thr Pro  
145 150 155 160

Val Ala Gly Asn Ser Val Ala Ser Ile Asp Ala Gln Leu Gly Asp Ala  
165 170 175

Arg Asp Ala Arg Ala Asp Ala Ala Ala Gln Leu Ser Lys Ala Gln Ser  
180 185 190

Gln Leu Asp Ala Thr Thr Val Leu Ser Thr Leu Glu Gly Thr Val Val  
195 200 205

Glu Val Asn Ser Asn Val Ser Lys Ser Pro Thr Gly Ala Ser Gln Val  
210 215 220

Met Val His Ile Val Ser Asn Glu Asn Leu Gln Val Lys Gly Glu Leu  
225 230 235 240

Ser Glu Tyr Asn Leu Ala Asn Leu Ser Val Gly Gln Glu Val Ser Phe  
245 250 255

Thr Ser Lys Val Tyr Pro Asp Lys Lys Trp Thr Gly Lys Leu Ser Tyr  
260 265 270

Ile Ser Asp Tyr Pro Lys Asn Asn Gly Glu Ala Ala Ser Pro Ala Ala  
275 280 285

Gly Asn Asn Thr Gly Ser Lys Tyr Pro Tyr Thr Ile Asp Val Thr Gly  
290 295 300

Glu Val Gly Asp Leu Lys Gln Gly Phe Ser Val Asn Ile Glu Val Lys  
 305 310 315 320

Ser Lys Thr Lys Ala Ile Leu Val Pro Val Ser Ser Leu Val Met Asp  
 325 330 335

Asp Ser Lys Asn Tyr Val Trp Ile Val Asp Glu Gln Gln Lys Ala Lys  
 340 345 350

Lys Val Glu Val Ser Leu Gly Asn Ala Asp Ala Glu Asn Gln Glu Ile  
 355 360 365

Thr Ser Gly Leu Thr Asn Gly Ala Lys Val Ile Ser Asn Pro Thr Ser  
 370 375 380

Ser Leu Glu Glu Gly Lys Glu Val Lys Ala Asp Glu Ala Thr Asn  
 385 390 395

<210> 95

<211> 759

<212> DNA

<213> Streptococcus pneumoniae

<400> 95

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759

&lt;210&gt; 96

&lt;211&gt; 252

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 96

Met Ser Arg Lys Pro Phe Ile Ala Gly Asn Trp Lys Met Asn Lys Asn  
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Pro Glu Glu Ala Lys Ala Phe Val Glu Ala Val Ala Ser Lys Leu Pro  
 20 25 30

Ser Ser Asp Leu Val Glu Ala Gly Ile Ala Ala Pro Ala Leu Asp Leu  
 35 40 45

Thr Thr Val Leu Ala Val Ala Lys Gly Ser Asn Leu Lys Val Ala Ala  
 50 55 60

Gln Asn Cys Tyr Phe Glu Asn Ala Gly Ala Phe Thr Gly Glu Thr Ser  
 65 70 75 80

Pro Gln Val Leu Lys Glu Ile Gly Thr Asp Tyr Val Val Ile Gly His  
 85 90 95

Ser Glu Arg Arg Asp Tyr Phe His Glu Thr Asp Glu Asp Ile Asn Lys  
 100 105 110

Lys Ala Lys Ala Ile Phe Ala Asn Gly Met Leu Pro Ile Ile Cys Cys  
 115 120 125

Gly Glu Ser Leu Glu Thr Tyr Glu Ala Gly Lys Ala Ala Glu Phe Val  
 130 135 140

Gly Ala Gln Val Ser Ala Ala Leu Ala Gly Leu Thr Ala Glu Gln Val  
 145 150 155 160

128

Ala Ala Ser Val Ile Ala Tyr Glu Pro Ile Trp Ala Ile Gly Thr Gly  
165 170 175

Lys Ser Ala Ser Gln Asp Asp Ala Gln Lys Met Cys Lys Val Val Arg  
180 185 190

Asp Val Val Ala Ala Asp Phe Gly Gln Glu Val Ala Asp Lys Val Arg  
195 200 205

Val Gln Tyr Gly Gly Ser Val Lys Pro Glu Asn Val Ala Ser Tyr Met  
210 215 220

Ala Cys Pro Asp Val Asp Gly Ala Leu Val Gly Gly Ala Ser Leu Glu  
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Ala Glu Ser Phe Leu Ala Leu Leu Asp Phe Val Lys  
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<210> 97

<211> 1473

<212> DNA

<213> Streptococcus pneumoniae

<400> 97

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<210> 98

<211> 490

<212> PRT

<213> Streptococcus pneumoniae

<400> 98

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 35 40 45

Lys Thr Gln Thr Ser Ala Glu Val Gln Thr Asn Ala Ala Ala His Trp  
 50 55 60

Asp Gly Asp Tyr Tyr Val Lys Asp Asp Gly Ser Lys Ala Gln Ser Glu  
 65 70 75 80

Trp Ile Phe Asp Asn Tyr Tyr Lys Ala Trp Phe Tyr Ile Asn Ser Asp  
 85 90 95

130

Gly Arg Tyr Ser Gln Asn Glu Trp His Gly Asn Tyr Tyr Leu Lys Ser  
100 105 110

Gly Gly Tyr Met Ala Gln Asn Glu Trp Ile Tyr Asp Ser Asn Tyr Lys  
115 120 125

Ser Trp Phe Tyr Leu Lys Ser Asp Gly Ala Tyr Ala His Gln Glu Trp  
130 135 140

Gln Leu Ile Gly Asn Lys Trp Tyr Tyr Phe Lys Lys Trp Gly Tyr Met  
145 150 155 160

Ala Lys Ser Gln Trp Gln Gly Ser Tyr Phe Leu Asn Gly Gln Gly Ala  
165 170 175

Met Met Gln Asn Glu Trp Leu Tyr Asp Pro Ala Tyr Ser Ala Tyr Phe  
180 185 190

Tyr Leu Lys Ser Asp Gly Thr Tyr Ala Asn Gln Glu Trp Gln Lys Val  
195 200 205

Gly Gly Lys Trp Tyr Tyr Phe Lys Lys Trp Gly Tyr Met Ala Arg Asn  
210 215 220

Glu Trp Gln Gly Asn Tyr Tyr Leu Thr Gly Ser Gly Ala Met Ala Thr  
225 230 235 240

Asp Glu Val Ile Met Asp Gly Thr Arg Tyr Ile Phe Ala Ala Ser Gly  
245 250 255

Glu Leu Lys Glu Lys Lys Asp Leu Asn Val Gly Trp Val His Arg Asp  
260 265 270

Gly Lys Arg Tyr Phe Phe Asn Asn Arg Glu Glu Gln Val Gly Thr Glu  
275 280 285

His Ala Lys Lys Val Ile Asp Ile Ser Glu His Asn Gly Arg Ile Asn  
290 295 300

131

Asp Trp Lys Lys Val Ile Asp Glu Asn Glu Val Asp Gly Val Ile Val  
305 310 315 320

Arg Leu Gly Tyr Ser Gly Lys Glu Asp Lys Glu Leu Ala His Asn Ile  
325 330 335

Lys Glu Leu Asn Arg Leu Gly Ile Pro Tyr Gly Val Tyr Leu Tyr Thr  
340 345 350

Tyr Ala Glu Asn Glu Thr Asp Ala Glu Ser Asp Ala Lys Gln Thr Ile  
355 360 365

Glu Leu Ile Lys Lys Tyr Asn Met Asn Leu Ser Tyr Pro Ile Tyr Tyr  
370 375 380

Asp Val Glu Asn Trp Glu Tyr Val Asn Lys Ser Lys Arg Ala Pro Ser  
385 390 395 400

Asp Thr Gly Thr Trp Val Lys Ile Ile Asn Lys Tyr Met Asp Thr Met  
405 410 415

Lys Gln Ala Gly Tyr Gln Asn Val Tyr Val Tyr Ser Tyr Arg Ser Leu  
420 425 430

Leu Gln Thr Arg Leu Lys His Pro Asp Ile Leu Lys His Val Asn Trp  
435 440 445

Val Ala Ala Tyr Thr Asn Ala Leu Glu Trp Glu Asn Pro His Tyr Ser  
450 455 460

Gly Lys Lys Gly Trp Gln Tyr Thr Ser Ser Glu Tyr Met Lys Gly Ile  
465 470 475 480

Gln Gly Arg Val Asp Val Ser Val Trp Tyr  
485 490



&lt;210&gt; 99

&lt;211&gt; 774

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 99

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&lt;210&gt; 100

&lt;211&gt; 257

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 100

Met Lys Lys Phe Ala Asn Leu Tyr Leu Gly Leu Val Phe Leu Val Leu

1

5

10

15

Tyr Leu Pro Ile Phe Tyr Leu Ile Gly Tyr Ala Phe Asn Ala Gly Asp

20

25

30

Asp Met Asn Ser Phe Thr Gly Phe Ser Trp Thr His Phe Glu Thr Met

35

40

45

Phe Gly Asp Gly Arg Leu Met Leu Ile Leu Ala Gln Thr Phe Phe Leu

50

55

60

133

Ala Phe Leu Ser Ala Leu Ile Ala Thr Ile Ile Gly Thr Phe Gly Ala  
65 70 75 80

Ile Tyr Ile Tyr Gln Ser Arg Lys Lys Tyr Gln Glu Ala Phe Leu Ser  
85 90 95

Leu Asn Asn Ile Leu Met Val Ala Pro Asp Val Met Ile Gly Ala Ser  
100 105 110

Phe Leu Ile Leu Phe Thr Gln Leu Lys Phe Ser Leu Gly Phe Leu Thr  
115 120 125

Val Leu Ser Ser His Val Ala Phe Ser Ile Pro Ile Val Val Leu Met  
130 135 140

Val Leu Pro Arg Leu Lys Glu Met Asn Gly Asp Met Ile His Ala Ala  
145 150 155 160

Tyr Asp Leu Gly Ala Ser Gln Phe Gln Met Phe Lys Glu Ile Met Leu  
165 170 175

Pro Tyr Leu Thr Pro Ser Ile Ile Thr Gly Tyr Phe Met Ala Phe Thr  
180 185 190

Tyr Ser Leu Asp Asp Phe Ala Val Thr Phe Phe Val Thr Gly Asn Gly  
195 200 205

Phe Ser Thr Leu Ser Val Glu Ile Tyr Ser Arg Ala Arg Lys Gly Ile  
210 215 220

Ser Leu Glu Ile Asn Ala Leu Ser Ala Leu Val Phe Leu Phe Ser Ile  
225 230 235 240

Ile Leu Val Val Gly Tyr Tyr Phe Ile Ser Arg Glu Lys Glu Glu Gln  
245 250 255

Ala

<210> 101

<211> 1071

<212> DNA

<213> Streptococcus pneumoniae

<400> 101

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<210> 102

<211> 356

<212> PRT

<213> Streptococcus pneumoniae

<400> 102

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1

5

10

15

135

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Asp Ser Gln Lys Leu Val Ile Tyr Asn Trp Gly Asp Tyr Ile Asp Pro  
35 40 45

Glu Leu Leu Thr Gln Phe Thr Glu Glu Thr Gly Ile Gln Val Gln Tyr  
50 55 60

Glu Thr Phe Asp Ser Asn Glu Ala Met Tyr Thr Lys Ile Lys Gln Gly  
65 70 75 80

Gly Thr Thr Tyr Asp Ile Ala Ile Pro Ser Glu Tyr Met Ile Asn Lys  
85 90 95

Met Lys Asp Glu Asp Leu Leu Val Pro Leu Asp Tyr Ser Lys Ile Glu  
100 105 110

Gly Ile Glu Asn Ile Gly Pro Glu Phe Leu Asn Gln Ser Phe Asp Pro  
115 120 125

Gly Asn Lys Phe Ser Ile Pro Tyr Phe Trp Gly Thr Leu Gly Ile Val  
130 135 140

Tyr Asn Glu Thr Met Val Asp Glu Ala Pro Glu His Trp Asp Asp Leu  
145 150 155 160

Trp Lys Pro Glu Tyr Lys Asn Ser Ile Met Leu Phe Asp Gly Ala Arg  
165 170 175

Glu Val Leu Gly Leu Gly Leu Asn Ser Leu Gly Tyr Ser Leu Asn Ser  
180 185 190

Lys Asp Leu Gln Gln Leu Glu Glu Thr Val Asp Lys Leu Tyr Lys Leu  
195 200 205

Thr Pro Asn Ile Lys Ala Ile Val Ala Asp Glu Met Lys Gly Tyr Met  
210 215 220

Variable	Mean	SD	Min	Max
Age	34.5	10.2	21	55
Gender	1.2	0.4	1	2
Marital status	1.8	0.4	1	2
Education	12.5	1.5	9	16
Income	1.5	0.5	1	2
Occupation	1.5	0.5	1	2
Health status	1.5	0.5	1	2
Stress level	1.5	0.5	1	2
Life satisfaction	1.5	0.5	1	2
Resilience	1.5	0.5	1	2
Optimism	1.5	0.5	1	2
Self-efficacy	1.5	0.5	1	2
Emotional stability	1.5	0.5	1	2
Neuroticism	1.5	0.5	1	2
Extraversion	1.5	0.5	1	2
Conscientiousness	1.5	0.5	1	2
Agreeableness	1.5	0.5	1	2
Openness	1.5	0.5	1	2
Depression	1.5	0.5	1	2
Anxiety	1.5	0.5	1	2
Stress management	1.5	0.5	1	2
Problem-solving	1.5	0.5	1	2
Emotional regulation	1.5	0.5	1	2
Resilience training	1.5	0.5	1	2
Optimism training	1.5	0.5	1	2
Self-efficacy training	1.5	0.5	1	2
Emotional stability training	1.5	0.5	1	2
Neuroticism training	1.5	0.5	1	2
Extraversion training	1.5	0.5	1	2
Conscientiousness training	1.5	0.5	1	2
Agreeableness training	1.5	0.5	1	2
Openness training	1.5	0.5	1	2
Depression training	1.5	0.5	1	2
Anxiety training	1.5	0.5	1	2
Stress management training	1.5	0.5	1	2
Problem-solving training	1.5	0.5	1	2
Emotional regulation training	1.5	0.5	1	2
Resilience training	1.5	0.5	1	2
Optimism training	1.5	0.5	1	2
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Openness training	1.5	0.5	1	2
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Anxiety training	1.5	0.5	1	2
Stress management training	1.5	0.5	1	2
Problem-solving training	1.5	0.5	1	2
Emotional regulation training	1.5	0.5	1	2
Resilience training	1.5	0.5	1	2
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Anxiety training	1.5	0.5	1	2
Stress management training	1.5	0.5	1	2
Problem-solving training	1.5	0.5	1	2
Emotional regulation training	1.5	0.5	1	2
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Optimism training	1.5	0.5	1	2
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Depression training	1.5	0.5	1	2
Anxiety training	1.5	0.5	1	2
Stress management training	1.5	0.5	1	2
Problem-solving training	1.5	0.5	1	2
Emotional regulation training	1.5	0.5	1	2
Resilience training	1.5	0.5	1	2
Optimism training	1.5	0.5	1	2
Self-efficacy training	1.5	0.5	1	2
Emotional stability training	1.5	0.5	1	2

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<210> 104

<211> 616

<212> PRT

<213> *Streptococcus pneumoniae*

<400> 104

Met Asn Lys Lys Leu Thr Asp Tyr Val Ile Asp Leu Val Glu Ile Leu

1

5

10

15

Asn Lys Gln Gln Lys Gln Val Phe Trp Gly Ile Phe Asp Ile Phe Ser  
                   20                  25                  30

Met Val Val Ser Ile Ile Val Ser Tyr Ile Leu Phe Tyr Gly Leu Ile  
                   35                  40                  45

Asn Pro Ala Pro Val Asp Tyr Ile Ile Tyr Thr Ser Leu Ala Phe Leu  
                   50                  55                  60

Phe Tyr Gln Leu Met Ile Gly Phe Trp Gly Leu Asn Ala Ser Ile Ser  
                   65                  70                  75                  80

Arg Tyr Ser Lys Ile Thr Asp Phe Met Lys Ile Phe Phe Gly Val Thr  
                   85                  90                  95

Ala Ser Ser Val Leu Ser Tyr Ser Ile Cys Tyr Ala Phe Leu Pro Leu  
                   100                  105                  110

Phe Ser Ile Arg Phe Ile Ile Leu Phe Ile Leu Leu Ser Thr Phe Leu  
                   115                  120                  125

Ile Leu Leu Pro Arg Ile Thr Trp Gln Leu Ile Tyr Ser Arg Arg Lys  
                   130                  135                  140

Lys Gly Ser Gly Asp Gly Glu His Arg Arg Thr Phe Leu Ile Gly Ala  
                   145                  150                  155                  160

Gly Asp Gly Gly Ala Leu Phe Met Asp Ser Tyr Gln His Pro Thr Ser  
                   165                  170                  175

Glu Leu Glu Leu Val Gly Ile Leu Asp Lys Asp Ser Lys Lys Lys Gly  
                   180                  185                  190

Gln Lys Leu Gly Gly Ile Pro Val Leu Gly Ser Tyr Asp Asn Leu Pro  
                   195                  200                  205

Glu Leu Ala Lys Arg His Gln Ile Glu Arg Val Ile Val Ala Ile Pro  
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139

Ser Leu Asp Pro Ser Glu Tyr Glu Arg Ile Leu Gln Met Cys Asn Lys  
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Leu Gly Val Lys Cys Tyr Lys Met Pro Lys Val Glu Thr Val Val Gln  
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Gly Leu His Gln Ala Gly Thr Gly Phe Gln Lys Ile Asp Ile Thr Asp  
260 265 270

Leu Leu Gly Arg Gln Glu Ile Arg Leu Asp Glu Ser Arg Leu Gly Ala  
275 280 285

Glu Leu Thr Gly Lys Thr Ile Leu Val Thr Gly Ala Gly Gly Ser Ile  
290 295 300

Gly Ser Glu Ile Cys Arg Gln Val Ser Arg Phe Asn Pro Glu Arg Ile  
305 310 315 320

Val Leu Leu Gly His Gly Glu Asn Ser Ile Tyr Leu Val Tyr His Glu  
325 330 335

Leu Ile Arg Lys Phe Gln Gly Ile Asp Tyr Val Pro Val Ile Ala Asp  
340 345 350

Ile Gln Asp Tyr Asp Arg Leu Leu Gln Val Phe Glu Gln Tyr Lys Pro  
355 360 365

Ala Ile Val Tyr His Ala Ala Ala His Lys His Val Pro Met Met Glu  
370 375 380

Arg Asn Pro Lys Glu Ala Phe Lys Asn Asn Ile Arg Gly Thr Tyr Asn  
385 390 395 400

Val Ala Lys Ala Val Asp Glu Ala Lys Val Ser Lys Met Val Met Ile  
405 410 415

Ser Thr Asp Lys Ala Val Asn Pro Pro Asn Val Met Gly Ala Thr Lys  
420 425 430



Arg Val Ala Glu Leu Ile Val Thr Gly Phe Asn Gln Arg Ser Gln Ser  
 435 440 445

Thr Tyr Cys Ala Val Arg Phe Gly Asn Val Leu Gly Ser Arg Gly Ser  
 450 455 460

Val Ile Pro Val Phe Glu Arg Gln Ile Ala Glu Gly Gly Pro Val Thr  
 465 470 475 480

Val Thr Asp Phe Arg Met Thr Arg Tyr Phe Met Thr Ile Pro Glu Ala  
 485 490 495

Ser Arg Leu Val Ile His Ala Gly Ala Tyr Ala Lys Asp Gly Glu Val  
 500 505 510

Phe Ile Leu Asp Met Gly Lys Pro Val Lys Ile Tyr Asp Leu Ala Lys  
 515 520 525

Lys Met Val Leu Leu Ser Gly His Thr Glu Ser Glu Ile Pro Ile Val  
 530 535 540

Glu Val Gly Ile Arg Pro Gly Glu Lys Leu Tyr Glu Glu Leu Leu Val  
 545 550 555 560

Ser Thr Glu Leu Val Asp Asn Gln Val Met Asp Lys Ile Phe Val Gly  
 565 570 575

Lys Val Asn Val Met Pro Leu Glu Ser Ile Asn Gln Lys Ile Gly Glu  
 580 585 590

Phe Arg Thr Leu Ser Gly Asp Glu Leu Lys Gln Ala Ile Ile Ala Phe  
 595 600 605

Ala Asn Gln Thr Thr His Ile Glu  
 610 615

&lt;210&gt; 105

&lt;211&gt; 1338

&lt;212&gt; DNA

<213> *Streptococcus pneumoniae*

&lt;400&gt; 105

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1338

&lt;210&gt; 106

&lt;211&gt; 445

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 106

Met Ile Glu Leu Tyr Asp Ser Tyr Ser Gln Glu Ser Arg Asp Leu His  
 1 5 10 15

Glu Ser Leu Val Ala Thr Gly Leu Ser Gln Leu Gly Val Val Ile Asp  
 20 25 30

Ala Asp Gly Phe Leu Pro Asp Gly Leu Leu Ser Pro Phe Thr Tyr Tyr  
 35 40 45

Leu Gly Tyr Glu Asp Gly Lys Pro Leu Tyr Phe Asn Gln Val Pro Val  
 50 55 60

Ser Asp Phe Trp Glu Ile Leu Gly Asp Asn Gln Ser Ala Cys Ile Glu  
 65 70 75 80

Asp Val Thr Gln Glu Arg Ala Val Ile His Tyr Ala Asp Gly Met Gln  
 85 90 95

Ala Arg Leu Val Lys Gln Val Asp Trp Lys Asp Leu Glu Gly Arg Val  
 100 105 110

Arg Gln Val Asp His Tyr Asn Arg Phe Gly Ala Cys Phe Ala Thr Thr  
 115 120 125

Thr Tyr Ser Ala Asp Ser Glu Pro Ile Met Thr Val Tyr Gln Asp Val  
 130 135 140

Asn Gly Gln Gln Val Leu Leu Glu Asn His Val Thr Gly Asp Ile Leu  
 145 150 155 160

Leu Thr Leu Pro Gly Gln Ser Met Arg Tyr Phe Ala Asn Lys Val Glu  
 165 170 175

Phe Ile Thr Phe Phe Leu Gln Asp Leu Glu Ile Asp Thr Ser Gln Leu  
 180 185 190

Ile Phe Asn Thr Leu Ala Thr Pro Phe Leu Val Ser Phe His His Pro  
 195 200 205

Asp Lys Ser Gly Ser Asp Val Leu Val Trp Gln Glu Pro Leu Tyr Asp  
 210 215 220

Ala Ile Pro Gly Asn Met Gln Leu Ile Leu Glu Ser Asp Asn Val Arg  
 225 230 235 240

Thr Lys Lys Ile Ile Ile Pro Asn Lys Ala Thr Tyr Glu Arg Ala Leu  
 245 250 255

Glu Leu Thr Asp Glu Lys Tyr His Asp Gln Phe Val His Leu Gly Tyr  
 260 265 270

His Tyr Gln Phe Lys Arg Asp Asn Phe Leu Arg Arg Asp Ala Leu Ile  
 275 280 285

Leu Thr Asn Ser Asp Gln Ile Glu Gln Val Glu Ala Ile Ala Gly Ala  
 290 295 300

Leu Pro Asp Val Thr Phe Arg Ile Ala Ala Val Thr Glu Met Ser Ser  
 305 310 315 320

Lys Leu Leu Asp Met Leu Cys Tyr Pro Asn Val Ala Leu Tyr Gln Asn  
 325 330 335

Ala Ser Pro Gln Lys Ile Gln Glu Leu Tyr Gln Leu Ser Asp Ile Tyr  
 340 345 350

Leu Asp Ile Asn His Ser Asn Glu Leu Leu Gln Ala Val Arg Gln Ala  
 355 360 365

Phe Glu His Asn Leu Leu Ile Leu Gly Phe Asn Gln Thr Val His Asn  
 370 375 380

Arg Leu Tyr Ile Ala Pro Asp His Leu Phe Glu Ser Ser Glu Val Ala  
 385 390 395 400

Ala Leu Val Glu Thr Ile Lys Leu Ala Leu Ser Asp Val Asp Gln Met  
 405 410 415

Arg Gln Ala Leu Gly Lys Gln Gly Gln His Ala Asn Tyr Val Asp Leu  
 420 425 430

Val Arg Tyr Gln Glu Thr Met Gln Thr Val Leu Gly Gly  
 435 440 445

<210> 107

<211> 1512

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 107

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 caggttatct ggctttataa tcatttcaca gatatcaaaa ttgcacctac tagcgtgaca 240  
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<210> 108

<211> 503

<212> PRT

<213> Streptococcus pneumoniae

<400> 108

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Leu Ser Ser Lys Phe Ile Phe Thr Asp Met Ile Leu Ala Asp Asn Ile  
 35 40 45

Gln His Leu Thr Ala Asn Ile Gly Phe Asp Asp Asn Gln Val Ile Trp  
 50 55 60

Leu Tyr Asn His Phe Thr Asp Ile Lys Ile Ala Pro Thr Ser Val Thr  
 65 70 75 80

Val Asp Asp Val Leu Ala Tyr Phe Gly Gly Glu Glu Ser His Arg Glu  
 85 90 95

Lys Asn Gly Lys Val Leu Arg Val Phe Phe Phe Asp Gln Asp Lys Phe  
 100 105 110

Val Thr Cys Tyr Leu Val Asp Glu Asn Lys Asp Leu Val Gln His Ala  
 115 120 125

Glu Tyr Val Phe Lys Gly Asn Leu Ile Arg Lys Asp Tyr Phe Ser Tyr  
 130 135 140

Thr Arg Tyr Cys Ser Glu Tyr Phe Ala Pro Lys Asp Asn Val Ala Val  
 145 150 155 160

Leu Tyr Gln Arg Thr Phe Tyr Asn Glu Asp Gly Thr Pro Val Tyr Asp  
 165 170 175

Ile Leu Met Asn Gln Gly Lys Glu Glu Val Tyr His Phe Lys Asp Lys  
 180 185 190

Ile Phe Tyr Gly Lys Gln Ala Phe Val Arg Ala Phe Met Lys Ser Leu  
 195 200 205

Asn Leu Asn Lys Ser Asp Leu Val Ile Leu Asp Arg Glu Thr Gly Ile  
 210 215 220

Gly Gln Val Val Phe Glu Glu Ala Gln Thr Ala His Leu Ala Val Val  
 225 230 235 240

Val His Ala Glu His Tyr Ser Glu Asn Ala Thr Asn Glu Asp Tyr Ile  
 245 250 255

Leu Trp Asn Asn Tyr Tyr Asp Tyr Gln Phe Thr Asn Ala Asp Lys Val  
 260 265 270

Asp Phe Phe Ile Val Ser Thr Asp Arg Gln Asn Glu Val Leu Gln Glu  
 275 280 285

Gln Phe Ala Lys Tyr Thr Gln His Gln Pro Lys Ile Val Thr Ile Pro  
 290 295 300

Val Gly Ser Ile Asp Ser Leu Thr Asp Ser Ser Gln Gly Arg Lys Pro  
 305 310 315 320

147

Phe Ser Leu Ile Thr Ala Ser Arg Leu Ala Lys Glu Lys His Ile Asp  
325 330 335

Trp Leu Val Lys Ala Val Ile Glu Ala His Lys Glu Leu Pro Glu Leu  
340 345 350

Thr Phe Asp Ile Tyr Gly Ser Gly Gly Glu Asp Ser Leu Leu Arg Glu  
355 360 365

Ile Ile Ala Asn His Gln Ala Glu Asp Tyr Ile Gln Leu Lys Gly His  
370 375 380

Ala Glu Leu Ser Gln Ile Tyr Ser Gln Tyr Glu Val Tyr Leu Thr Ala  
385 390 395 400

Ser Thr Ser Glu Gly Phe Gly Leu Thr Leu Met Glu Ala Ile Gly Ser  
405 410 415

Gly Leu Pro Leu Ile Gly Phe Asp Val Pro Tyr Gly Asn Gln Thr Phe  
420 425 430

Ile Glu Asp Gly Gln Asn Gly Tyr Leu Ile Pro Ser Ser Ser Asp His  
435 440 445

Val Glu Asp Gln Ile Lys Gln Ala Tyr Ala Ala Lys Ile Cys Gln Leu  
450 455 460

Tyr Gln Glu Asn Arg Leu Glu Ala Met Arg Ala Tyr Ser Tyr Gln Ile  
465 470 475 480

Ala Glu Gly Phe Leu Thr Lys Glu Ile Leu Glu Lys Trp Lys Lys Thr  
485 490 495

Val Glu Glu Val Leu His Asp  
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&lt;210&gt; 109

&lt;211&gt; 2292

&lt;212&gt; DNA

<213> *Streptococcus pneumoniae*

&lt;400&gt; 109

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<210> 110

<211> 763

<212> PRT

<213> Streptococcus pneumoniae

<400> 110

Met Ser Ser Leu Ser Asp Gln Glu Leu Val Ala Lys Thr Val Glu Phe  
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Arg Gln Arg Leu Ser Glu Gly Glu Ser Leu Asp Asp Ile Leu Val Glu  
 20 25 30

Ala Phe Ala Val Val Arg Glu Ala Asp Lys Arg Ile Leu Gly Met Phe  
 35 40 45

Pro Tyr Asp Val Gln Val Met Gly Ala Ile Val Met His Tyr Gly Asn  
 50 55 60

Val Ala Glu Met Asn Thr Gly Glu Gly Lys Thr Leu Thr Ala Thr Met  
 65 70 75 80

Pro Val Tyr Leu Asn Ala Phe Ser Gly Glu Gly Val Met Val Val Thr  
 85 90 95

Pro Asn Glu Tyr Leu Ser Lys Arg Asp Ala Glu Glu Met Gly Gln Val  
 100 105 110

Tyr Arg Phe Leu Gly Leu Thr Ile Gly Val Pro Phe Thr Glu Asp Pro  
 115 120 125

150

Lys Lys Glu Met Lys Ala Glu Glu Lys Lys Leu Ile Tyr Ala Ser Asp  
130 135 140

Ile Ile Tyr Thr Thr Asn Ser Asn Leu Gly Phe Asp Tyr Leu Asn Asp  
145 150 155 160

Asn Leu Ala Ser Asn Glu Glu Gly Lys Phe Leu Arg Pro Phe Asn Tyr  
165 170 175

Val Ile Ile Asp Glu Ile Asp Asp Ile Leu Leu Asp Ser Ala Gln Thr  
180 185 190

Pro Leu Ile Ile Ala Gly Ser Pro Arg Val Gln Ser Asn Tyr Tyr Ala  
195 200 205

Ile Ile Asp Thr Leu Val Thr Thr Leu Val Glu Gly Glu Asp Tyr Ile  
210 215 220

Phe Lys Glu Glu Lys Glu Glu Val Trp Leu Thr Thr Lys Gly Ala Lys  
225 230 235 240

Ser Ala Glu Asn Phe Leu Gly Ile Asp Asn Leu Tyr Lys Glu Glu His  
245 250 255

Ala Ser Phe Ala Arg His Leu Val Tyr Ala Ile Arg Ala His Lys Leu  
260 265 270

Phe Thr Lys Asp Lys Asp Tyr Ile Ile Arg Gly Asn Glu Met Val Leu  
275 280 285

Val Asp Lys Gly Thr Gly Arg Leu Met Glu Met Thr Lys Leu Gln Gly  
290 295 300

Gly Leu His Gln Ala Ile Glu Ala Lys Glu His Val Lys Leu Ser Pro  
305 310 315 320

Glu Thr Arg Ala Met Ala Ser Ile Thr Tyr Gln Ser Leu Phe Lys Met  
325 330 335

Phe Asn Lys Ile Ser Gly Met Thr Gly Thr Gly Lys Val Ala Glu Lys  
 340 345 350

Glu Phe Ile Glu Thr Tyr Asn Met Ser Val Val Arg Ile Pro Thr Asn  
 355 360 365

Arg Pro Arg Gln Arg Ile Asp Tyr Pro Asp Asn Leu Tyr Ile Thr Leu  
 370 375 380

Pro Glu Lys Val Tyr Ala Ser Leu Glu Tyr Ile Lys Gln Tyr His Ala  
 385 390 395 400

Lys Gly Asn Pro Leu Leu Val Phe Val Gly Ser Val Glu Met Ser Gln  
 405 410 415

Leu Tyr Ser Ser Leu Leu Phe Arg Glu Gly Ile Ala His Asn Val Leu  
 420 425 430

Asn Ala Asn Asn Ala Ala Arg Glu Ala Gln Ile Ile Ser Glu Ser Gly  
 435 440 445

Gln Met Gly Ala Val Thr Val Ala Thr Ser Met Ala Gly Arg Gly Thr  
 450 455 460

Asp Ile Lys Leu Gly Lys Gly Val Ala Glu Leu Gly Gly Leu Ile Val  
 465 470 475 480

Ile Gly Thr Glu Arg Met Glu Ser Gln Arg Ile Asp Leu Gln Ile Arg  
 485 490 495

Gly Arg Ser Gly Arg Gln Gly Asp Pro Gly Met Ser Lys Phe Phe Val  
 500 505 510

Ser Leu Glu Asp Asp Val Ile Lys Lys Phe Gly Pro Ser Trp Val His  
 515 520 525

Lys Lys Tyr Lys Asp Tyr Gln Val Gln Asp Met Thr Gln Pro Glu Val  
 530 535 540

152

Leu Lys Gly Arg Lys Tyr Arg Lys Leu Val Glu Lys Ala Gln His Ala  
545 550 555 560

Ser Asp Ser Ala Gly Arg Ser Ala Arg Arg Gln Thr Leu Glu Tyr Ala  
565 570 575

Glu Ser Met Asn Ile Gln Arg Asp Ile Val Tyr Lys Glu Arg Asn Arg  
580 585 590

Leu Ile Asp Gly Ser Arg Asp Leu Glu Asp Val Val Val Asp Ile Ile  
595 600 605

Glu Arg Tyr Thr Glu Glu Val Ala Ala Asp His Tyr Ala Ser Arg Glu  
610 615 620

Leu Leu Phe His Phe Ile Val Thr Asn Ile Ser Phe His Val Lys Glu  
625 630 635 640

Val Pro Asp Tyr Ile Asp Val Thr Asp Lys Thr Ala Val Arg Ser Phe  
645 650 655

Met Lys Gln Val Ile Asp Lys Glu Leu Ser Glu Lys Lys Glu Leu Leu  
660 665 670

Asn Gln His Asp Leu Tyr Glu Gln Phe Leu Arg Leu Ser Leu Leu Lys  
675 680 685

Ala Ile Asp Asp Asn Trp Val Glu Gln Val Asp Tyr Leu Gln Gln Leu  
690 695 700

Ser Met Ala Ile Gly Gly Gln Ser Ala Ser Gln Lys Asn Pro Ile Val  
705 710 715 720

Glu Tyr Tyr Gln Glu Ala Tyr Ala Gly Phe Glu Ala Met Lys Glu Gln  
725 730 735

Ile His Ala Asp Met Val Arg Asn Leu Leu Met Gly Leu Val Glu Val  
740 745 750

Thr Pro Lys Gly Glu Ile Val Thr His Phe Pro

755

760

&lt;210&gt; 111

&lt;211&gt; 879

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 111

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&lt;210&gt; 112

&lt;211&gt; 292

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 112

Met Lys Gln Glu Trp Phe Glu Ser Asn Asp Phe Val Lys Thr Thr Ser

1

5

10

15

Lys Asn Lys Pro Glu Glu Gln Ala Gln Glu Val Ala Asp Lys Ala Glu  
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Glu Arg Ile Pro Asp Leu Asp Thr Pro Ile Glu Lys Asn Thr Gln Leu  
 35 40 45

Glu Glu Glu Val Ser Gln Ala Glu Val Glu Leu Glu Ser Gln Gln Glu  
 50 55 60

Glu Lys Ile Glu Ala Pro Glu Asp Ser Glu Ala Arg Thr Glu Ile Glu  
 65 70 75 80

Glu Lys Lys Ala Ser Asn Ser Thr Glu Glu Glu Pro Asp Leu Ser Lys  
 85 90 95

Glu Thr Glu Lys Val Thr Ile Ala Glu Glu Ser Gln Glu Ala Leu Pro  
 100 105 110

Gln Gln Lys Ala Thr Thr Lys Glu Pro Leu Leu Ile Ser Lys Ser Leu  
 115 120 125

Glu Ser Pro Tyr Ile Pro Asp Gln Ala Pro Lys Ser Arg Asp Lys Trp  
 130 135 140

Lys Glu Gln Val Leu Asp Phe Trp Ser Trp Leu Val Glu Ala Ile Lys  
 145 150 155 160

Ser Pro Thr Ser Lys Leu Glu Thr Ser Ile Thr His Ser Tyr Thr Ala  
 165 170 175

Phe Leu Leu Leu Ile Leu Phe Ser Ala Ser Ser Phe Phe Phe Ser Ile  
 180 185 190

Tyr His Ile Lys His Ala Tyr Tyr Gly His Ile Ala Ser Ile Asn Ser  
 195 200 205

Arg Phe Pro Glu Gln Leu Ala Pro Leu Thr Leu Phe Ser Ile Ile Ser  
 210 215 220

155

Ile Leu Val Ala Thr Thr Leu Phe Phe Phe Ser Phe Leu Leu Gly Ser  
225 230 235 240

Phe Val Val Arg Arg Phe Ile His Gln Glu Lys Asp Trp Thr Leu Asp  
245 250 255

Lys Val Leu Gln Gln Tyr Ser Gln Leu Leu Ala Ile Pro Ile Ser Ser  
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Leu Leu Leu Leu Val Ser Leu Leu Ser Leu Ile Ala Tyr Asp Leu Gln  
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Pro Ser Cys Val  
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<210> 113

<211> 327

<212> DNA

<213> Streptococcus pneumoniae

<400> 113

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<210> 114

<211> 108

<212> PRT

<213> Streptococcus pneumoniae

<400> 114

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 20 25 30

Lys Leu Ile Ala Asn Ile Lys Glu Ser Thr Leu Tyr Pro Ile Leu Lys  
 35 40 45

Lys Leu Glu Gly Asn Ser Phe Leu Thr Thr Tyr Ser Arg Glu Phe Gln  
 50 55 60

Gly Arg Met Arg Lys Tyr Tyr Ser Leu Thr Asn Gly Gly Ile Glu Gln  
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Gly Ile Ile Glu Gly Ser Ile Arg His Asp Lys Asn  
 100 105

<210> 115

<211> 954

<212> DNA

<213> Streptococcus pneumoniae

<400> 115

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<210> 116

<211> 317

<212> PRT

<213> Streptococcus pneumoniae

<400> 116

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Val Gln Val Ile Gln Ser Asp Leu Ala Thr Asn Phe Tyr Asp Ala Leu

20

25

30

Val Glu Gln Asn Ser Ile Tyr Leu Asp Gly Glu Thr Glu Leu Asn Gln

35

40

45

Val Lys Asp Asn Asn Gln Ala Leu Lys Arg Leu Ala Leu Arg Lys Glu

50

55

60

Glu Trp Leu Lys Thr Tyr Gln Phe Leu Leu Met Lys Ala Gly Gln Thr

65

70

75

80

Glu Pro Leu Gln Ala Asn His Gln Phe Thr Pro Asp Ala Ile Ala Leu

85

90

95

Leu Leu Val Phe Ile Val Glu Glu Leu Phe Lys Glu Glu Glu Ile Thr

100

105

110

Ile Leu Glu Met Gly Ser Gly Met Gly Ile Leu Gly Ala Ile Phe Leu

115

120

125

Thr Ser Leu Thr Lys Lys Val Asp Tyr Leu Gly Met Glu Val Asp Asp

130

135

140

158

Leu Leu Ile Asp Leu Ala Ala Ser Met Ala Asp Val Ile Gly Leu Gln  
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Ala Gly Phe Val Gln Gly Asp Ala Val Arg Pro Gln Met Leu Lys Glu  
165 170 175

Ser Asp Val Val Ile Ser Asp Leu Pro Val Gly Tyr Tyr Pro Asp Asp  
180 185 190

Ala Val Ala Ser Arg His Gln Val Ala Ser Ser Gln Glu His Thr Tyr  
195 200 205

Ala His His Leu Leu Met Glu Gln Gly Leu Lys Tyr Leu Lys Ser Asp  
210 215 220

Gly Tyr Ala Ile Phe Leu Ala Pro Ser Asp Leu Leu Thr Ser Pro Gln  
225 230 235 240

Ser Asp Leu Leu Lys Glu Trp Leu Lys Glu Glu Ala Ser Leu Val Ala  
245 250 255

Met Ile Ser Leu Pro Glu Asn Leu Phe Ala Asn Ala Lys Gln Ser Lys  
260 265 270

Thr Ile Phe Ile Leu Gln Lys Lys Asn Glu Ile Ala Val Glu Pro Phe  
275 280 285

Val Tyr Pro Leu Ala Ser Leu Gln Asp Ala Ser Val Leu Met Lys Phe  
290 295 300

Lys Glu Asn Phe Gln Lys Trp Thr Gln Gly Thr Glu Ile  
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&lt;210&gt; 117

&lt;211&gt; 1902

&lt;212&gt; DNA

<213> *Streptococcus pneumoniae*

&lt;400&gt; 117

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&lt;210&gt; 118

&lt;211&gt; 633

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 118

Met Ile Ile Leu Gln Ala Asn Lys Ile Glu Arg Ser Phe Ala Gly Glu

1

5

10

15

Val Leu Phe Asp Asn Ile Asn Leu Gln Val Asp Glu Arg Asp Arg Ile

20

25

30

Ala Leu Val Gly Lys Asn Gly Ala Gly Lys Ser Thr Leu Leu Lys Ile

35

40

45

Leu Val Gly Glu Glu Glu Pro Thr Ser Gly Glu Ile Asn Lys Lys Lys

50

55

60

Asp Ile Ser Leu Ser Tyr Leu Ala Gln Asp Ser Arg Phe Glu Ser Glu

65

70

75

80

Asn Thr Ile Tyr Asp Glu Met Leu His Val Phe Asn Asp Leu Arg Arg

85

90

95

Thr Glu Arg Gln Leu Arg Gln Met Glu Leu Glu Met Gly Glu Lys Ser

100

105

110

Gly Glu Asp Leu Asp Lys Leu Met Ser Asp Tyr Asp Arg Leu Ser Glu

115

120

125

Asn Phe Arg Gln Ala Gly Gly Phe Thr Tyr Glu Ala Asp Ile Arg Ala

130

135

140

Ile Leu Asn Gly Phe Lys Phe Asp Glu Ser Met Trp Gln Met Lys Ile

145

150

155

160

161

Ala Glu Leu Ser Gly Gly Gln Asn Thr Arg Leu Ala Leu Ala Lys Met  
165 170 175

Leu Leu Glu Lys Pro Asn Leu Leu Val Leu Asp Glu Pro Thr Asn His  
180 185 190

Leu Asp Ile Glu Thr Ile Ala Trp Leu Glu Asn Tyr Leu Val Asn Tyr  
195 200 205

Ser Gly Ala Leu Ile Ile Val Ser His Asp Arg Tyr Phe Leu Asp Lys  
210 215 220

Val Ala Thr Ile Thr Leu Asp Leu Thr Lys His Ser Leu Asp Arg Tyr  
225 230 235 240

Val Gly Asn Tyr Ser Arg Phe Val Glu Leu Lys Glu Gln Lys Leu Val  
245 250 255

Thr Glu Ala Lys Asn Tyr Glu Lys Gln Gln Lys Glu Ile Ala Ala Leu  
260 265 270

Glu Asp Phe Val Asn Arg Asn Leu Val Arg Ala Ser Thr Thr Lys Arg  
275 280 285

Ala Gln Ser Arg Arg Lys Gln Leu Glu Lys Met Glu Arg Leu Asp Lys  
290 295 300

Pro Glu Ala Gly Lys Lys Ala Ala Asn Met Thr Phe Gln Ser Glu Lys  
305 310 315 320

Thr Ser Gly Asn Val Val Leu Thr Val Glu Asn Ala Ala Val Gly Tyr  
325 330 335

Asp Gly Glu Val Leu Ser Gln Pro Ile Asn Leu Asp Leu Arg Lys Met  
340 345 350

Asn Ala Val Ala Ile Val Gly Pro Asn Gly Ile Gly Lys Ser Thr Phe  
355 360 365

Ile Lys Ser Ile Val Asp Gln Ile Pro Phe Ile Lys Gly Glu Lys Arg  
 370 375 380

Phe Gly Ala Asn Val Glu Val Gly Tyr Tyr Asp Gln Thr Gln Ser Lys  
 385 390 395 400

Leu Thr Pro Ser Asn Thr Val Leu Asp Glu Leu Trp Asn Asp Phe Lys  
 405 410 415

Leu Thr Pro Glu Val Glu Ile Arg Asn Arg Leu Gly Ala Phe Leu Phe  
 420 425 430

Ser Gly Asp Asp Val Lys Lys Ser Val Gly Met Leu Ser Gly Gly Glu  
 435 440 445

Lys Ala Arg Leu Leu Leu Ala Lys Leu Ser Met Glu Asn Asn Asn Phe  
 450 455 460

Leu Ile Leu Asp Glu Pro Thr Asn His Leu Asp Ile Asp Ser Lys Glu  
 465 470 475 480

Val Leu Glu Asn Ala Leu Ile Asp Phe Asp Gly Thr Leu Leu Phe Val  
 485 490 495

Ser His Asp Arg Tyr Phe Ile Asn Arg Val Ala Thr His Val Leu Glu  
 500 505 510

Leu Ser Glu Asn Gly Ser Thr Leu Tyr Leu Gly Asp Tyr Asp Tyr Tyr  
 515 520 525

Val Glu Lys Lys Ala Thr Ala Glu Met Ser Gln Thr Glu Glu Ala Ser  
 530 535 540

Thr Ser Asn Gln Ala Lys Glu Ala Ser Pro Val Asn Asp Tyr Gln Ala  
 545 550 555 560

Gln Lys Glu Ser Gln Lys Glu Val Arg Lys Leu Met Arg Gln Ile Glu  
 565 570 575

Ser Leu Glu Ala Glu Ile Glu Glu Leu Glu Ser Gln Ser Gln Ala Ile  
 580 585 590

Ser Glu Gln Met Leu Glu Thr Asn Asp Ala Asp Lys Leu Met Glu Leu  
 595 600 605

Gln Ala Glu Leu Asp Lys Ile Ser His Arg Gln Glu Glu Ala Met Leu  
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Glu Trp Glu Glu Leu Ser Glu Gln Val  
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<210> 119

<211> 1179

<212> DNA

<213> Streptococcus pneumoniae

<400> 119

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<210> 120

<211> 392

<212> PRT

<213> Streptococcus pneumoniae

<400> 120

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Met Gly Thr Ile Gly Gln Thr Val Leu Gly Met Tyr Gln Ile Ser Glu  
 35 40 45

Leu Val Thr Ser Ile Leu Val Asn Pro Phe Gly Gly Val Ile Ser Asp  
 50 55 60

Arg Phe Ser Arg Arg Lys Ile Leu Met Thr Ala Asp Leu Val Cys Gly  
 65 70 75 80

Ile Leu Cys Leu Ala Ile Ser Phe Ile Arg Asn Asp Ser Trp Met Ile  
 85 90 95

Gly Ala Leu Ile Val Ala Asn Ile Val Gln Ala Ile Ala Phe Ala Phe  
 100 105 110

Ser Arg Thr Ala Asn Lys Ala Ile Ile Thr Glu Val Val Glu Lys Asp  
 115 120 125

Glu Ile Val Ile Tyr Asn Ser Arg Leu Glu Leu Val Leu Gln Val Val  
 130 135 140

165

Gly Val Ser Ser Pro Val Leu Ser Phe Leu Val Leu Gln Phe Ala Ser  
145 150 155 160

Leu His Met Thr Leu Leu Leu Asp Ser Leu Thr Phe Phe Ile Ala Phe  
165 170 175

Val Leu Val Ala Phe Leu Pro Lys Glu Glu Ala Lys Val Gln Glu Lys  
180 185 190

Lys Ala Phe Thr Gly Arg Asp Ile Phe Val Asp Ile Lys Asp Gly Leu  
195 200 205

His Tyr Ile Trp His Gln Gln Glu Ile Phe Phe Leu Leu Leu Val Ala  
210 215 220

Ser Ser Val Asn Phe Phe Phe Ala Ala Phe Glu Phe Leu Leu Pro Phe  
225 230 235 240

Ser Asn Gln Leu Tyr Gly Ser Glu Gly Ala Tyr Ala Ser Ile Leu Thr  
245 250 255

Met Gly Ala Ile Gly Ser Ile Ile Gly Ala Leu Leu Ala Ser Lys Ile  
260 265 270

Lys Ala Asn Ile Tyr Asn Leu Leu Ile Leu Leu Ala Leu Thr Gly Val  
275 280 285

Gly Val Phe Met Met Gly Leu Pro Leu Pro Thr Phe Leu Ser Phe Ser  
290 295 300

Gly Asn Leu Val Cys Glu Leu Phe Met Thr Ile Phe Asn Ile His Phe  
305 310 315 320

Phe Thr Gln Val Gln Thr Lys Val Glu Ser Glu Phe Leu Gly Arg Val  
325 330 335

Leu Ser Thr Ile Phe Thr Leu Ala Ile Leu Phe Met Pro Ile Ala Lys  
340 345 350

Gly Phe Met Thr Val Leu Pro Ser Val His Leu Tyr Ser Phe Leu Ile  
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Arg Thr His Phe Glu Lys Leu Ile  
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<210> 121

<211> 2466

<212> DNA

<213> Streptococcus pneumoniae

<400> 121

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<210> 122

<211> 821

<212> PRT

<213> Streptococcus pneumoniae

<400> 122

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10

15

Gln Gln Lys Gln Lys Asn Lys Lys Ser Ala Arg Pro Gly Lys Lys Gly

20

25

30

Ser Ser Thr Lys Lys Ser Lys Thr Leu Asp Lys Ser Ala Ile Phe Pro

35

40

45

Ala Ile Leu Leu Ser Ile Lys Ala Leu Phe Asn Leu Leu Phe Val Leu  
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Gly Phe Leu Gly Gly Met Leu Gly Ala Gly Ile Ala Leu Gly Tyr Gly  
 65 70 75 80

Val Ala Leu Phe Asp Lys Val Arg Val Pro Gln Thr Glu Glu Leu Val  
 85 90 95

Asn Gln Val Lys Asp Ile Ser Ser Ile Ser Glu Ile Thr Tyr Ser Asp  
 100 105 110

Gly Thr Val Ile Ala Ser Ile Glu Ser Asp Leu Leu Arg Thr Ser Ile  
 115 120 125

Ser Ser Glu Gln Ile Ser Glu Asn Leu Lys Lys Ala Ile Ile Ala Thr  
 130 135 140

Glu Asp Glu His Phe Lys Glu His Lys Gly Val Val Pro Lys Ala Val  
 145 150 155 160

Ile Arg Ala Thr Leu Gly Lys Phe Val Gly Leu Gly Ser Ser Ser Gly  
 165 170 175

Gly Ser Thr Leu Thr Gln Gln Leu Ile Lys Gln Gln Val Val Gly Asp  
 180 185 190

Ala Pro Thr Leu Ala Arg Lys Ala Ala Glu Ile Val Asp Ala Leu Ala  
 195 200 205

Leu Glu Arg Ala Met Asn Lys Asp Glu Ile Leu Thr Thr Tyr Leu Asn  
 210 215 220

Val Ala Pro Phe Gly Arg Asn Asn Lys Gly Gln Asn Ile Ala Gly Ala  
 225 230 235 240

Arg Gln Ala Ala Glu Gly Ile Phe Gly Val Asp Ala Ser Gln Leu Thr  
 245 250 255

Val Pro Gln Ala Ala Phe Leu Ala Gly Leu Pro Gln Ser Pro Ile Thr  
 260 265 270

Tyr Ser Pro Tyr Glu Asn Thr Gly Glu Leu Lys Ser Asp Glu Asp Leu  
 275 280 285

Glu Ile Gly Leu Arg Arg Ala Lys Ala Val Leu Tyr Ser Met Tyr Arg  
 290 295 300

Thr Gly Ala Leu Ser Lys Asp Glu Tyr Ser Gln Tyr Lys Asp Tyr Asp  
 305 310 315 320

Leu Lys Gln Asp Phe Leu Pro Ser Gly Thr Val Thr Gly Ile Ser Arg  
 325 330 335

Asp Tyr Leu Tyr Phe Thr Thr Leu Ala Glu Ala Gln Glu Arg Met Tyr  
 340 345 350

Asp Tyr Leu Ala Gln Arg Asp Asn Val Ser Ala Lys Glu Leu Lys Asn  
 355 360 365

Glu Ala Thr Gln Lys Phe Tyr Arg Asp Leu Ala Ala Lys Glu Ile Glu  
 370 375 380

Asn Gly Gly Tyr Lys Ile Thr Thr Thr Ile Asp Gln Lys Ile His Ser  
 385 390 395 400

Ala Met Gln Ser Ala Val Ala Asp Tyr Gly Tyr Leu Leu Asp Asp Gly  
 405 410 415

Thr Gly Arg Val Glu Val Gly Asn Val Leu Met Asp Asn Gln Thr Gly  
 420 425 430

Ala Ile Leu Gly Phe Val Gly Gly Arg Asn Tyr Gln Glu Asn Gln Asn  
 435 440 445

Asn His Ala Phe Asp Thr Lys Arg Ser Pro Ala Ser Thr Thr Lys Pro  
 450 455 460

170

Leu Leu Ala Tyr Gly Ile Ala Ile Asp Gln Gly Leu Met Gly Ser Glu  
465 470 475 480

Thr Ile Leu Ser Asn Tyr Pro Thr Asn Phe Ala Asn Gly Asn Pro Ile  
485 490 495

Met Tyr Ala Asn Ser Lys Gly Thr Gly Met Met Thr Leu Gly Glu Ala  
500 505 510

Leu Asn Tyr Ser Trp Asn Ile Pro Ala Tyr Trp Thr Tyr Arg Met Leu  
515 520 525

Arg Glu Lys Gly Val Asp Val Lys Gly Tyr Met Glu Lys Met Gly Tyr  
530 535 540

Glu Ile Pro Glu Tyr Gly Ile Glu Ser Leu Pro Met Gly Gly Gly Ile  
545 550 555 560

Glu Val Thr Val Ala Gln His Thr Asn Gly Tyr Gln Thr Leu Ala Asn  
565 570 575

Asn Gly Val Tyr His Gln Lys His Val Ile Ser Lys Ile Glu Ala Ala  
580 585 590

Asp Gly Arg Val Val Tyr Glu Tyr Gln Asp Lys Pro Val Gln Val Tyr  
595 600 605

Ser Lys Ala Thr Ala Thr Ile Met Gln Gly Leu Leu Arg Glu Val Leu  
610 615 620

Ser Ser Arg Val Thr Thr Thr Phe Lys Ser Asn Leu Thr Ser Leu Asn  
625 630 635 640

Pro Thr Leu Ala Asn Ala Asp Trp Ile Gly Lys Thr Gly Thr Thr Asn  
645 650 655

Gln Asp Glu Asn Met Trp Leu Met Leu Ser Thr Pro Arg Leu Thr Leu  
660 665 670

171

Gly Gly Trp Ile Gly His Asp Asp Asn His Ser Leu Ser Arg Arg Ala  
675 680 685

Gly Tyr Ser Asn Asn Ser Asn Tyr Met Ala His Leu Val Asn Ala Ile  
690 695 700

Gln Gln Ala Ser Pro Ser Ile Trp Gly Asn Glu Arg Phe Ala Leu Asp  
705 710 715 720

Pro Ser Val Val Lys Ser Glu Val Leu Lys Ser Thr Gly Gln Lys Pro  
725 730 735

Glu Lys Val Ser Val Glu Gly Lys Glu Val Glu Val Thr Gly Ser Thr  
740 745 750

Val Thr Ser Tyr Trp Ala Asn Lys Ser Gly Ala Pro Ala Thr Ser Tyr  
755 760 765

Arg Phe Ala Ile Gly Gly Ser Asp Ala Asp Tyr Gln Asn Ala Trp Ser  
770 775 780

Ser Ile Val Gly Ser Leu Pro Thr Pro Ser Ser Ser Ser Ser Ser  
785 790 795 800

Ser Ser Ser Ser Asp Ser Ser Asn Ser Ser Thr Thr Arg Pro Ser Ser  
805 810 815

Ser Arg Ala Arg Arg  
820

<210> 123

<211> 1974

<212> DNA

<213> Streptococcus pneumoniae

<400> 123

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gtcttatcca ctttcattat ttttgtaat aataatctgt tgacggtttt aattttgttt 120



cttttttagtag gaggotatgt ttttttattt aagaaactga gagtgcatta tacaaggagt 180  
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 gaacagatgc ctgtaggtgt tatgaaattg aatttatott ctggagaggt tgagtggttt 300  
 aatccctatg ctgaattgat tttagaccaag gaagatggtg attttgattt agaagctggt 360  
 caaacgatta tcaaggcttc agtaggaaat ccgtctactt atgccaaagct tggtgagaag 420  
 cgttatgctg ttcatatgga tgcttcttcc ggtgttttgt atttttaga tgtatccagg 480  
 gaacaagcca taacagatga attggtaaca agtagaccag tgattgggat tgtctctgtg 540  
 gataattatg atgatttgga ggatgaaact tctgagtcag atattagtca aatcaatagt 600  
 tttgtagcta attttatatc agagttttca gaaaaacaca tgatgttttc tcgtcgggta 660  
 agtatggatc gatttttatct atttactgac tacacggtgc ttgagggctt gatgaatgat 720  
 aaattttctg ttattgatgc ttccagagaa gagtcgaaac agagacagtt gcccttgacc 780  
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 tcagctcgaa gtcgtagtaa actgaatgta caacggatta tggaagagtt aggcggtgga 1860  
 ggccacttta atttggcagc agctcaaatt aaagatgtaa ccttgtcaga agcaggtgaa 1920  
 aaactgacag aaattgtatt aaatgaaatg aaggaaaagg agaaaagaaga atga 1974

<210> 124

<211> 657

<212> PRT

<213> *Streptococcus pneumoniae*

Met Lys Lys Phe Tyr Val Ser Pro Ile Phe Pro Ile Leu Val Gly Leu  
1 5 10 15

Leu Leu Thr Val Leu Ile Leu Phe Leu Phe Val Gly Gly Tyr Val Phe  
35 40 45

Ile Gln Tyr Val Asn His Gln Ala Glu Glu Ser Leu Thr Ala Leu Leu  
65 70 75 80

Val Glu Trp Phe Asn Pro Tyr Ala Glu Leu Ile Leu Thr Lys Glu Asp  
100 105 110

Gly Asn Pro Ser Thr Tyr Ala Lys Leu Gly Glu Lys Arg Tyr Ala Val  
130 135 140

Glu Gln Ala Ile Thr Asp Glu Leu Val Thr Ser Arg Pro Val Ile Gly  
165 170 175

Ile Val Ser Val Asp Asn Tyr Asp Asp Leu Glu Asp Glu Thr Ser Glu  
180 185 190

Ser Asp Ile Ser Gln Ile Asn Ser Phe Val Ala Asn Phe Ile Ser Glu  
 195 200 205

Phe Ser Glu Lys His Met Met Phe Ser Arg Arg Val Ser Met Asp Arg  
 210 215 220

Phe Tyr Leu Phe Thr Asp Tyr Thr Val Leu Glu Gly Leu Met Asn Asp  
 225 230 235 240

Lys Phe Ser Val Ile Asp Ala Phe Arg Glu Glu Ser Lys Gln Arg Gln  
 245 250 255

Leu Pro Leu Thr Leu Ser Met Gly Phe Ser Tyr Gly Asp Gly Asn His  
 260 265 270

Asp Glu Ile Gly Lys Val Ala Leu Leu Asn Leu Asn Leu Ala Glu Val  
 275 280 285

Arg Gly Gly Asp Gln Val Val Val Lys Glu Asn Asp Glu Thr Lys Asn  
 290 295 300

Pro Val Tyr Phe Gly Gly Gly Ser Ala Ala Ser Ile Lys Arg Thr Arg  
 305 310 315 320

Thr Arg Thr Arg Ala Met Met Thr Ala Ile Ser Asp Lys Ile Arg Ser  
 325 330 335

Val Asp Gln Val Phe Val Val Gly His Lys Asn Leu Asp Met Asp Ala  
 340 345 350

Leu Gly Ser Ala Val Gly Met Gln Leu Phe Ala Ser Asn Val Ile Glu  
 355 360 365

Asn Ser Tyr Ala Leu Tyr Asp Glu Glu Gln Met Ser Pro Asp Ile Glu  
 370 375 380

Arg Ala Val Ser Phe Ile Glu Lys Glu Gly Val Thr Lys Leu Leu Ser  
 385 390 395 400

175

Val Lys Asp Ala Met Gly Met Val Thr Asn Arg Ser Leu Leu Ile Leu  
405 410 415

Val Asp His Ser Lys Thr Ala Leu Thr Leu Ser Lys Glu Phe Tyr Asp  
420 425 430

Leu Phe Thr Gln Thr Ile Val Ile Asp His His Arg Arg Asp Gln Asp  
435 440 445

Phe Pro Asp Asn Ala Val Ile Thr Tyr Ile Glu Ser Gly Ala Ser Ser  
450 455 460

Ala Ser Glu Leu Val Thr Glu Leu Ile Gln Phe Gln Asn Ser Lys Lys  
465 470 475 480

Asn Arg Leu Ser Arg Met Gln Ala Ser Val Leu Met Ala Gly Met Met  
485 490 495

Leu Asp Thr Lys Asn Phe Thr Ser Arg Val Thr Ser Arg Thr Phe Asp  
500 505 510

Val Ala Ser Tyr Leu Arg Thr Arg Gly Ser Asp Ser Ile Ala Ile Gln  
515 520 525

Glu Ile Ala Ala Thr Asp Phe Glu Glu Tyr Arg Glu Val Asn Glu Leu  
530 535 540

Ile Leu Gln Gly Arg Lys Leu Gly Ser Asp Val Leu Ile Ala Glu Ala  
545 550 555 560

Lys Asp Met Lys Cys Tyr Asp Thr Val Val Ile Ser Lys Ala Ala Asp  
565 570 575

Ala Met Leu Ala Met Ser Gly Ile Glu Ala Ser Phe Val Leu Ala Lys  
580 585 590

Asn Thr Gln Gly Phe Ile Ser Ile Ser Ala Arg Ser Arg Ser Lys Leu  
595 600 605

Asn Val Gln Arg Ile Met Glu Glu Leu Gly Gly Gly Gly His Phe Asn  
 610 615 620

Leu Ala Ala Ala Gln Ile Lys Asp Val Thr Leu Ser Glu Ala Gly Glu  
 625 630 635 640

Lys Leu Thr Glu Ile Val Leu Asn Glu Met Lys Glu Lys Glu Lys Glu  
 645 650 655

Glu

<210> 125

<211> 663

<212> DNA

<213> Streptococcus pneumoniae

<400> 125

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 ggggaagaga actgtccaaa ttgtatgaaa acagagttgt caacaaagtg tcaagattgt 180  
 caactttggt gtaaagaggg agttgaagtc agtcatagag cgatTTTTac ttacaatcaa 240  
 gctatgaagg atTTTTcag tcggtataag tttgatggag acttctctgtt aagaaaagtt 300  
 ttgcgttcat ttttaagtga ggagttgaaa aagtacaaag agtatcaatt tgttgtaatt 360  
 cccctaagtc ctgatagata tgctaataga ggatttaatac aggttgaggg cttggttagag 420  
 gcagcaggct ttgagtatct ggatttatta gagaaaagag aagagagagc cagttcttct 480  
 aaaaatcggt cagagcgctt ggggacagaa cttcctttct ttattaaaag tggagtcact 540  
 attcctaaaa aaatcctact tatagatgat atctatacta caggagcaac tataaatcgt 600  
 gttaagaaac tgttgaaga agctggtgct aaggatgtaa aaacattttc ccttgtaaga 660  
 tga 663

<210> 126

<211> 220

<212> PRT

<213> Streptococcus pneumoniae

&lt;400&gt; 126

Met Lys Cys Leu Leu Cys Gly Gln Thr Met Lys Thr Val Leu Thr Phe  
 1 5 10 15

Ser Ser Leu Leu Leu Leu Arg Asn Asp Asp Ser Cys Leu Cys Ser Asp  
 20 25 30

Cys Asp Ser Thr Phe Glu Arg Ile Gly Glu Glu Asn Cys Pro Asn Cys  
 35 40 45

Met Lys Thr Glu Leu Ser Thr Lys Cys Gln Asp Cys Gln Leu Trp Cys  
 50 55 60

Lys Glu Gly Val Glu Val Ser His Arg Ala Ile Phe Thr Tyr Asn Gln  
 65 70 75 80

Ala Met Lys Asp Phe Phe Ser Arg Tyr Lys Phe Asp Gly Asp Phe Leu  
 85 90 95

Leu Arg Lys Val Phe Ala Ser Phe Leu Ser Glu Glu Leu Lys Lys Tyr  
 100 105 110

Lys Glu Tyr Gln Phe Val Val Ile Pro Leu Ser Pro Asp Arg Tyr Ala  
 115 120 125

Asn Arg Gly Phe Asn Gln Val Glu Gly Leu Val Glu Ala Ala Gly Phe  
 130 135 140

Glu Tyr Leu Asp Leu Leu Glu Lys Arg Glu Glu Arg Ala Ser Ser Ser  
 145 150 155 160

Lys Asn Arg Ser Glu Arg Leu Gly Thr Glu Leu Pro Phe Phe Ile Lys  
 165 170 175

Ser Gly Val Thr Ile Pro Lys Lys Ile Leu Leu Ile Asp Asp Ile Tyr  
 180 185 190

Thr Thr Gly Ala Thr Ile Asn Arg Val Lys Lys Leu Leu Glu Glu Ala  
 195 200 205

Gly Ala Lys Asp Val Lys Thr Phe Ser Leu Val Arg  
 210 215 220

<210> 127

<211> 1299

<212> DNA

<213> Streptococcus pneumoniae

<400> 127

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atgaaagtaa atttagatta tctcggtcgt ttattttactg agaatgaatt aacagaagaa 60
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caacgctgta atagtactat tctagaagaa tggatatttc ccatcgggtgc ttactattgt 180
cgagagtgtc tgctgatgaa gcgagtcaga agtgatcaaa ctttatacta ttttccgcag 240
gaggattttc caaagcaaga tggttctcaa tggcgcggcc aattaactcc ttttcaagag 300
aaggtgtcag agggattgtc tcaagtagta gacaagcaaa agccaacctt agttcatgcy 360
gtaacaggag ctggaaagac agaaatgatt tatcaagtag tggctaaagt gatcaatgcy 420
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cgaacaccac tagttgttgc aacaacccat cagttattga agttttatca agcttttgat 600
ttgctgatag tggatgaagt agatgctttt cttatgttg ataatcccat gctttaccac 660
gctgtcaaga atagtgtaaa ggagaatgga ttgagaatct ttttaacagc gacttcgacc 720
aatgagttag ataaaaaggt ccgttttaga gaactaaaaa gactgaattt accgagacgg 780
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ttagacaaga atcgtttgtc accaaagtta aagtcctata ttgagaagca gagaaagaca 900
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ccgacaggag atttgctttt cttccatgat gggttaaatg cttcaatcaa gaaggcgatt 1260
aaggaaattc agatgatgaa taaggaggct ggtctatga 1299

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&lt;210&gt; 128

&lt;211&gt; 432

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 128

Met Lys Val Asn Leu Asp Tyr Leu Gly Arg Leu Phe Thr Glu Asn Glu

1

5

10

15

Leu Thr Glu Glu Glu Arg Gln Leu Ala Glu Lys Leu Pro Ala Met Arg

20

25

30

Lys Glu Lys Gly Lys Leu Phe Cys Gln Arg Cys Asn Ser Thr Ile Leu

35

40

45

Glu Glu Trp Tyr Leu Pro Ile Gly Ala Tyr Tyr Cys Arg Glu Cys Leu

50

55

60

Leu Met Lys Arg Val Arg Ser Asp Gln Thr Leu Tyr Tyr Phe Pro Gln

65

70

75

80

Glu Asp Phe Pro Lys Gln Asp Val Leu Lys Trp Arg Gly Gln Leu Thr

85

90

95

Pro Phe Gln Glu Lys Val Ser Glu Gly Leu Leu Gln Val Val Asp Lys

100

105

110

Gln Lys Pro Thr Leu Val His Ala Val Thr Gly Ala Gly Lys Thr Glu

115

120

125

Met Ile Tyr Gln Val Val Ala Lys Val Ile Asn Ala Gly Gly Ala Val

130

135

140

Cys Leu Ala Ser Pro Arg Ile Asp Val Cys Leu Glu Leu Tyr Lys Arg

145

150

155

160

Leu Gln Gln Asp Phe Ser Cys Gly Ile Ala Leu Leu His Gly Glu Ser

165

170

175



180

Glu Pro Tyr Phe Arg Thr Pro Leu Val Val Ala Thr Thr His Gln Leu  
180 185 190

Leu Lys Phe Tyr Gln Ala Phe Asp Leu Leu Ile Val Asp Glu Val Asp  
195 200 205

Ala Phe Pro Tyr Val Asp Asn Pro Met Leu Tyr His Ala Val Lys Asn  
210 215 220

Ser Val Lys Glu Asn Gly Leu Arg Ile Phe Leu Thr Ala Thr Ser Thr  
225 230 235 240

Asn Glu Leu Asp Lys Lys Val Arg Leu Gly Glu Leu Lys Arg Leu Asn  
245 250 255

Leu Pro Arg Arg Phe His Gly Asn Pro Leu Ile Ile Pro Lys Pro Ile  
260 265 270

Trp Leu Ser Asp Phe Asn Arg Tyr Leu Asp Lys Asn Arg Leu Ser Pro  
275 280 285

Lys Leu Lys Ser Tyr Ile Glu Lys Gln Arg Lys Thr Ala Tyr Pro Leu  
290 295 300

Leu Ile Phe Ala Ser Glu Ile Lys Lys Gly Glu Gln Leu Ala Glu Ile  
305 310 315 320

Leu Gln Glu Gln Phe Pro Asn Glu Lys Ile Gly Phe Val Ser Ser Val  
325 330 335

Thr Glu Asp Arg Leu Glu Gln Val Gln Ala Phe Arg Asp Gly Glu Leu  
340 345 350

Thr Ile Leu Ile Ser Thr Thr Ile Leu Glu Arg Gly Val Thr Phe Pro  
355 360 365

Cys Val Asp Val Phe Val Val Glu Ala Asn His Arg Leu Phe Thr Lys  
370 375 380

Ser Ser Leu Ile Gln Ile Gly Gly Arg Val Gly Arg Ser Met Asp Arg  
 385 390 395 400

Pro Thr Gly Asp Leu Leu Phe Phe His Asp Gly Leu Asn Ala Ser Ile  
 405 410 415

Lys Lys Ala Ile Lys Glu Ile Gln Met Met Asn Lys Glu Ala Gly Leu  
 420 425 430

<210> 129

<211> 870

<212> DNA

<213> Streptococcus pneumoniae

<400> 129

atgcaaattc aaaaaagttt taaggggcag tctccctatg gcaagctgta tctagtggca 60  
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 gactggattg ctgctgagga tacgcgcaat acagggtttt tgctcaagca ttttgacatt 180  
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 aaaaaagatt atcctgaaac acagattttt tatgaatcac ctcatcgtgt agcagacacg 540  
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 aaggacgagg aagacttggt cgtagaaatt caaacccgca tccagcaagg tgtgaagaaa 780  
 aaccaagcta tcaaggaagt cgctaagatt taccagtgga ataaaagtca gctctacgct 840  
 gcctaccacg actggaaga aaaacaataa 870

&lt;210&gt; 130

&lt;211&gt; 289

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 130

Met Gln Ile Gln Lys Ser Phe Lys Gly Gln Ser Pro Tyr Gly Lys Leu

1

5

10

15

Tyr Leu Val Ala Thr Pro Ile Gly Asn Leu Asp Asp Met Thr Phe Arg

20

25

30

Ala Ile Gln Thr Leu Lys Glu Val Asp Trp Ile Ala Ala Glu Asp Thr

35

40

45

Arg Asn Thr Gly Leu Leu Leu Lys His Phe Asp Ile Ser Thr Lys Gln

50

55

60

Ile Ser Phe His Glu His Asn Ala Lys Glu Lys Ile Pro Asp Leu Ile

65

70

75

80

Gly Phe Leu Lys Ala Gly Gln Ser Ile Ala Gln Val Ser Asp Ala Gly

85

90

95

Leu Pro Ser Ile Ser Asp Pro Gly His Asp Leu Val Lys Ala Ala Ile

100

105

110

Glu Glu Glu Ile Ala Val Val Thr Val Pro Gly Ala Ser Ala Gly Ile

115

120

125

Ser Ala Leu Ile Ala Ser Gly Leu Ala Pro Gln Pro His Ile Phe Tyr

130

135

140

Gly Phe Leu Pro Arg Lys Ser Gly Gln Gln Lys Gln Phe Phe Gly Leu

145

150

155

160

Lys Lys Asp Tyr Pro Glu Thr Gln Ile Phe Tyr Glu Ser Pro His Arg

165

170

175

183

Val Ala Asp Thr Leu Glu Asn Met Leu Glu Val Tyr Gly Asp Arg Ser  
180 185 190

Val Val Leu Val Arg Glu Leu Thr Lys Ile Tyr Glu Glu Tyr Gln Arg  
195 200 205

Gly Thr Ile Ser Glu Leu Leu Glu Ser Ile Ala Glu Thr Pro Leu Lys  
210 215 220

Gly Glu Cys Leu Leu Ile Val Glu Gly Ala Ser Gln Gly Val Glu Glu  
225 230 235 240

Lys Asp Glu Glu Asp Leu Phe Val Glu Ile Gln Thr Arg Ile Gln Gln  
245 250 255

Gly Val Lys Lys Asn Gln Ala Ile Lys Glu Val Ala Lys Ile Tyr Gln  
260 265 270

Trp Asn Lys Ser Gln Leu Tyr Ala Ala Tyr His Asp Trp Glu Glu Lys  
275 280 285

Gln

<210> 131

<211> 345

<212> DNA

<213> Streptococcus pneumoniae

<400> 131

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aagagcctgg tagaggaaaa tacagctctt cgcttggaag atagtaagtt gcgagaacgc 180  
ttgggtgagg tggaagcaga tgctcctgtc aaggccaagc atgttcgcga aagtgtccgt 240  
cgtattttacc gtgatggatt tcacgtatgt aatgattttt atggacaacg tcgagagcag 300  
gacgaagaat gtatgttttg tgacgagttg ttatacaggg agtaa 345

<210> 132

<211> 114

<212> PRT

<213> Streptococcus pneumoniae

<400> 132

Met Ile Lys Lys Gly Lys Gly Cys Phe Met Asp Lys Lys Glu Leu Phe

1

5

10

15

Asp Ala Leu Asp Asp Phe Ser Gln Gln Leu Leu Val Thr Leu Ala Asp

20

25

30

Val Glu Ala Ile Lys Lys Asn Leu Lys Ser Leu Val Glu Glu Asn Thr

35

40

45

Ala Leu Arg Leu Glu Asn Ser Lys Leu Arg Glu Arg Leu Gly Glu Val

50

55

60

Glu Ala Asp Ala Pro Val Lys Ala Lys His Val Arg Glu Ser Val Arg

65

70

75

80

Arg Ile Tyr Arg Asp Gly Phe His Val Cys Asn Asp Phe Tyr Gly Gln

85

90

95

Arg Arg Glu Gln Asp Glu Glu Cys Met Phe Cys Asp Glu Leu Leu Tyr

100

105

110

Arg Glu

<210> 133

<211> 639

<212> DNA

<213> Streptococcus pneumoniae

&lt;400&gt; 133

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cctggcggag tcttgattgg ggagaagatt cgggaagtga ttttggatcc aagtcatact 180
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gaaaaagttc ttccagccct tgaagctggc aagtgtgtca tcatggatcg ttttatcgat 300
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aatcagtttg cgacagatgg cctcaaaccg gatttgacac tctattttga catcgagggtg 420
gaagaagggc tggctcgtat tgctgctaag agtgaccgag aggttaatcg tttggatttg 480
gaagggttgg acttgcataa aaaagttcgt caaggctacc tttctcttct ggataaagag 540
ggaaatcgca ttgtcaagat tgatgctagt ctccctttgg agcaagttgt ggaaactacc 600
aaggctgtct tgtttgacgg aatgggcttg gccaaatga                               639

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&lt;210&gt; 134

&lt;211&gt; 212

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 134

```

Met Ser Lys Gly Phe Leu Val Ser Leu Glu Gly Pro Glu Gly Ala Gly
  1              5              10             15

```

```

Lys Thr Ser Val Leu Glu Ala Leu Leu Pro Ile Leu Glu Glu Lys Gly
      20              25              30

```

```

Val Glu Val Leu Thr Thr Arg Glu Pro Gly Gly Val Leu Ile Gly Glu
      35              40              45

```

```

Lys Ile Arg Glu Val Ile Leu Asp Pro Ser His Thr Gln Met Asp Ala
      50              55              60

```

```

Lys Thr Glu Leu Leu Leu Tyr Ile Ala Ser Arg Arg Gln His Leu Val
      65              70              75             80

```

```

Glu Lys Val Leu Pro Ala Leu Glu Ala Gly Lys Leu Val Ile Met Asp
      85              90              95

```

186

Arg Phe Ile Asp Ser Ser Val Ala Tyr Gln Gly Phe Gly Arg Gly Leu  
100 105 110

Asp Ile Glu Ala Ile Asp Trp Leu Asn Gln Phe Ala Thr Asp Gly Leu  
115 120 125

Lys Pro Asp Leu Thr Leu Tyr Phe Asp Ile Glu Val Glu Glu Gly Leu  
130 135 140

Ala Arg Ile Ala Ala Asn Ser Asp Arg Glu Val Asn Arg Leu Asp Leu  
145 150 155 160

Glu Gly Leu Asp Leu His Lys Lys Val Arg Gln Gly Tyr Leu Ser Leu  
165 170 175

Leu Asp Lys Glu Gly Asn Arg Ile Val Lys Ile Asp Ala Ser Leu Pro  
180 185 190

Leu Glu Gln Val Val Glu Thr Thr Lys Ala Val Leu Phe Asp Gly Met  
195 200 205

Gly Leu Ala Lys  
210

<210> 135

<211> 474

<212> DNA

<213> Streptococcus pneumoniae

<400> 135

atggtagaac aaagaaaatc aattaccatg aaagatggtg ctttagaagc aggagttagt 60  
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aaagtggaac aagcgattaa aactttgaat tacattccag attactacgc tagaggaatg 180  
aaaaaaaaatc gaacagaaac gattgcaatc attgtaccaa gtatctggca tccottcttt 240  
tcagaatttg ctatgcatgt ggaaaatgaa gtctataaga gaaataacaa attactctta 300  
tggttctatca atggtacaaa tagagagcaa gactatctgg agatggtgcg tcataataaa 360

gttgatggag tggttgccat tacctatagg ccaattgaac attacttgac gtcaggaatt 420  
 ccctttgtta gtattgaccg cacataactca gagattgcca ttccttgtgt ttca 474

<210> 136

<211> 158

<212> PRT

<213> Streptococcus pneumoniae

<400> 136

Met Val Glu Gln Arg Lys Ser Ile Thr Met Lys Asp Val Ala Leu Glu  
 1 5 10 15

Ala Gly Val Ser Val Gly Thr Val Ser Arg Val Ile Asn Lys Glu Lys  
 20 25 30

Gly Ile Lys Glu Val Thr Leu Lys Lys Val Glu Gln Ala Ile Lys Thr  
 35 40 45

Leu Asn Tyr Ile Pro Asp Tyr Tyr Ala Arg Gly Met Lys Lys Asn Arg  
 50 55 60

Thr Glu Thr Ile Ala Ile Ile Val Pro Ser Ile Trp His Pro Phe Phe  
 65 70 75 80

Ser Glu Phe Ala Met His Val Glu Asn Glu Val Tyr Lys Arg Asn Asn  
 85 90 95

Lys Leu Leu Leu Cys Ser Ile Asn Gly Thr Asn Arg Glu Gln Asp Tyr  
 100 105 110

Leu Glu Met Leu Arg His Asn Lys Val Asp Gly Val Val Ala Ile Thr  
 115 120 125

Tyr Arg Pro Ile Glu His Tyr Leu Thr Ser Gly Ile Pro Phe Val Ser  
 130 135 140



Ile Asp Arg Thr Tyr Ser Glu Ile Ala Ile Pro Cys Val Ser  
 145 150 155

<210> 137

<211> 374

<212> DNA

<213> Streptococcus pneumoniae

<400> 137

atgaatatat ttagaacaaa gaatgttagt ttagataaaa cagagatgca taggcatttg 60  
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 atcacaggta ctgcagctgc aacacttgct ggcccagccc tagtgatttc aatcgttatt 180  
 tctgccttgt gtgtgggatt atcagccctc ttttttgcag aatttgcctc gcgagtaccc 240  
 gctacaggag gtgcctatag ttacctctat gctatcttag gagaattccc tgcttggttg 300  
 gctggttggt taaccatgat ggagttcatg acagccatat caggcgtagc ttcgggttg 360  
 gcagcttatt ttaa 374

<210> 138

<211> 124

<212> PRT

<213> Streptococcus pneumoniae

<400> 138

Met Asn Ile Phe Arg Thr Lys Asn Val Ser Leu Asp Lys Thr Glu Met  
 1 5 10 15

His Arg His Leu Lys Leu Trp Asp Leu Ile Leu Leu Gly Ile Gly Ala  
 20 25 30

Met Val Gly Thr Gly Val Phe Thr Ile Thr Gly Thr Ala Ala Ala Thr  
 35 40 45

Leu Ala Gly Pro Ala Leu Val Ile Ser Ile Val Ile Ser Ala Leu Cys  
 50 55 60

Val Gly Leu Ser Ala Leu Phe Phe Ala Glu Phe Ala Ser Arg Val Pro  
 65 70 75 80

Ala Thr Gly Gly Ala Tyr Ser Tyr Leu Tyr Ala Ile Leu Gly Glu Phe  
 85 90 95

Pro Ala Trp Leu Ala Gly Trp Leu Thr Met Met Glu Phe Met Thr Ala  
 100 105 110

Ile Ser Gly Val Ala Ser Gly Trp Ala Ala Tyr Phe  
 115 120

<210> 139

<211> 1311

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 139

atgaaatcaa gagtaaagga aacgagtatg gataaaattg tggttcaagg tggcgataat 60  
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 gcgactattc tagcaagtga aggaaagacc gtcttgcaga atgttccgat tttgtcggat 180  
 gtctttatta tgaatcaggt agttggtggt ttgaatgcc aagttgactt tgatgaggaa 240  
 gctcatcttg tcaaggtgga tgctactggc gacatcactg aggaagcccc ttacaagtat 300  
 gtcagcaaga tgcgcgcctc catcgttgta ttagggccaa tccttgcccg tgtgggtcat 360  
 gccaaggtat ccatgccagg tggttgtacg attggtagcc gtcctattga tcttcatttg 420  
 aaaggtcttg aagctatggg ggttaagatt agtcagacag ctggttacat cgaagccaag 480  
 gcagaacgct tgcattggtg tcatatctat atggactttc caagtgttgg tgcaacgcag 540  
 aacttgatga tggcagcgac tctggctgat ggggtgacag tgattgagaa tgctgcgcgt 600  
 gagcctgaga ttgttgactt agccattctc cttaatgaaa tgggagccaa ggtcaaaggt 660  
 gctggtacag agactataac cattactggt gttgagaaac ttcattggtac gactcacaat 720  
 gtagtccaag accgtatcga agcaggaacc tttatggtag ctgctgccat gactgggtgt 780  
 gatgtcttga ttogagacgc tgtctgggag cacaaccgtc ccttgattgc caagttactt 840  
 gaaatgggtg ttgaagtaat tgaagaagac gaaggaattc gtgttcgttc tcaactagaa 900  
 aatctaaaag ctgttcatgt gaaaaccttg cccacccag gatttccaac agatatgcag 960  
 gctcaattta cagccttgat gacagttgca aaaggcgaat caaccatggt ggagacagtt 1020  
 ttcgaaaatc gtttccaaca cctagaagag atgcgcgcga tgggcttgca ttctgagatt 1080

atccgtgata cagctcgtat tgttggtgga cagcctttgc agggagcaga agttctttca 1140  
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 gtggtcggta aattggttca cttggataga ggttactacg gtttccatga gaagttggcg 1260  
 cagctagggtg ctaagattca gcggattgag gcaagtgatg aagatgaata a 1311

<210> 140

<211> 436

<212> PRT

<213> Streptococcus pneumoniae

<400> 140

Met Lys Ser Arg Val Lys Glu Thr Ser Met Asp Lys Ile Val Val Gln

1 5 10 15

Gly Gly Asp Asn Arg Leu Val Gly Ser Val Thr Ile Glu Gly Ala Lys

20 25 30

Asn Ala Val Leu Pro Leu Leu Ala Ala Thr Ile Leu Ala Ser Glu Gly

35 40 45

Lys Thr Val Leu Gln Asn Val Pro Ile Leu Ser Asp Val Phe Ile Met

50 55 60

Asn Gln Val Val Gly Gly Leu Asn Ala Lys Val Asp Phe Asp Glu Glu

65 70 75 80

Ala His Leu Val Lys Val Asp Ala Thr Gly Asp Ile Thr Glu Glu Ala

85 90 95

Pro Tyr Lys Tyr Val Ser Lys Met Arg Ala Ser Ile Val Val Leu Gly

100 105 110

Pro Ile Leu Ala Arg Val Gly His Ala Lys Val Ser Met Pro Gly Gly

115 120 125

Cys Thr Ile Gly Ser Arg Pro Ile Asp Leu His Leu Lys Gly Leu Glu

130 135 140

191

Ala Met Gly Val Lys Ile Ser Gln Thr Ala Gly Tyr Ile Glu Ala Lys  
145 150 155 160

Ala Glu Arg Leu His Gly Ala His Ile Tyr Met Asp Phe Pro Ser Val  
165 170 175

Gly Ala Thr Gln Asn Leu Met Met Ala Ala Thr Leu Ala Asp Gly Val  
180 185 190

Thr Val Ile Glu Asn Ala Ala Arg Glu Pro Glu Ile Val Asp Leu Ala  
195 200 205

Ile Leu Leu Asn Glu Met Gly Ala Lys Val Lys Gly Ala Gly Thr Glu  
210 215 220

Thr Ile Thr Ile Thr Gly Val Glu Lys Leu His Gly Thr Thr His Asn  
225 230 235 240

Val Val Gln Asp Arg Ile Glu Ala Gly Thr Phe Met Val Ala Ala Ala  
245 250 255

Met Thr Gly Gly Asp Val Leu Ile Arg Asp Ala Val Trp Glu His Asn  
260 265 270

Arg Pro Leu Ile Ala Lys Leu Leu Glu Met Gly Val Glu Val Ile Glu  
275 280 285

Glu Asp Glu Gly Ile Arg Val Arg Ser Gln Leu Glu Asn Leu Lys Ala  
290 295 300

Val His Val Lys Thr Leu Pro His Pro Gly Phe Pro Thr Asp Met Gln  
305 310 315 320

Ala Gln Phe Thr Ala Leu Met Thr Val Ala Lys Gly Glu Ser Thr Met  
325 330 335

Val Glu Thr Val Phe Glu Asn Arg Phe Gln His Leu Glu Glu Met Arg  
340 345 350

Arg Met Gly Leu His Ser Glu Ile Ile Arg Asp Thr Ala Arg Ile Val  
 355 360 365

Gly Gly Gln Pro Leu Gln Gly Ala Glu Val Leu Ser Thr Asp Leu Arg  
 370 375 380

Ala Ser Ala Ala Leu Ile Leu Thr Gly Leu Val Ala Gln Gly Glu Thr  
 385 390 395 400

Val Val Gly Lys Leu Val His Leu Asp Arg Gly Tyr Tyr Gly Phe His  
 405 410 415

Glu Lys Leu Ala Gln Leu Gly Ala Lys Ile Gln Arg Ile Glu Ala Ser  
 420 425 430

Asp Glu Asp Glu  
 435

<210> 141

<211> 1101

<212> DNA

<213> Streptococcus pneumoniae

<400> 141

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 aacaaaacaa gttataccgt acagtatggt gatactttga gcaccattgc agaagccttg 180  
 ggtgtagatg tcacagtgct tgcgaatctg aacaaaatca ctaatatgga cttgattttc 240  
 ccagaaactg ttttgacaac gactgtcaat gaagcagaag aagtaacaga agttgaaatc 300  
 caaacacctc aagcagactc tagtgaagaa gtgacaactg cgacagcaga tttgaccact 360  
 aatcaagtga ccgttgatga tcaaactgtt caggttgcag acctttctca accaattgca 420  
 gaagttacaa agacagtgat tgcttctgaa gaagtggcac catctacggg cacttctgtc 480  
 ccagaggagc aaacgaccga aacaactcgc ccagttgcag aagaagctcc tcaggaaacg 540  
 actccagctg agaagcagga aacacaaaca agccctcaag ctgcatcagc agtggaagca 600  
 actacaacaa gttcagaagc aaaagaagta gcatcatcaa atggagctac agcagcagtt 660  
 tctacttatt aaccagaaga aacgaaagta atttcaacaa cttacgaggc tccagctgcg 720

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cccgattatg ctggacttgc agtagcaaaa tctgaaaatg caggtcttca accacaaaca 780
gctgccttta agaagaaatt gctaacttgt ttggcattac atcctttagt ggttatcgtc 840
caggagacag tggagatcac ggaaaagggt tggctatcga ctttatggta ccagaacgtt 900
cagaattagg ggataagatt gcggaatatg ctattcaaaa tatggccagc cgtggcatta 960
gttacatcat ctggaaacaa cgtttctatg ctccattcga tagcaaatat gggccagcta 1020
acaattggaa cccaatgcc aaccgtggta gtgtgacaga aaatcactat gatcacgttc 1080
acgtttcaat gaatggataa                                     1100

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<210> 142

<211> 302

<212> PRT

<213> Streptococcus pneumoniae

<400> 142

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Met Leu Leu Ala Ser Thr Val Ala Leu Ser Phe Ala Pro Val Leu Ala
  1              5              10             15

Thr Gln Ala Glu Glu Val Leu Trp Thr Ala Arg Ser Val Glu Gln Ile
          20              25             30

Gln Asn Asp Leu Thr Lys Thr Asp Asn Lys Thr Ser Tyr Thr Val Gln
          35              40             45

Tyr Gly Asp Thr Leu Ser Thr Ile Ala Glu Ala Leu Gly Val Asp Val
          50              55             60

Thr Val Leu Ala Asn Leu Asn Lys Ile Thr Asn Met Asp Leu Ile Phe
          65              70             75             80

Pro Glu Thr Val Leu Thr Thr Thr Val Asn Glu Ala Glu Glu Val Thr
          85              90             95

Glu Val Glu Ile Gln Thr Pro Gln Ala Asp Ser Ser Glu Glu Val Thr
          100             105            110

Thr Ala Thr Ala Asp Leu Thr Thr Asn Gln Val Thr Val Asp Asp Gln
          115             120            125

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194

Thr Val Gln Val Ala Asp Leu Ser Gln Pro Ile Ala Glu Val Thr Lys  
130 135 140

Thr Val Ile Ala Ser Glu Glu Val Ala Pro Ser Thr Gly Thr Ser Val  
145 150 155 160

Pro Glu Glu Gln Thr Thr Glu Thr Thr Arg Pro Val Ala Glu Glu Ala  
165 170 175

Pro Gln Glu Thr Thr Pro Ala Glu Lys Gln Glu Thr Gln Thr Ser Pro  
180 185 190

Gln Ala Ala Ser Ala Val Glu Ala Thr Thr Thr Ser Ser Glu Ala Lys  
195 200 205

Glu Val Ala Ser Ser Asn Gly Ala Thr Ala Ala Val Ser Thr Tyr Gln  
210 215 220

Pro Glu Glu Thr Lys Val Ile Ser Thr Thr Tyr Glu Ala Pro Ala Ala  
225 230 235 240

Pro Asp Tyr Ala Gly Leu Ala Val Ala Lys Ser Glu Asn Ala Gly Leu  
245 250 255

Gln Pro Gln Thr Ala Ala Phe Lys Lys Lys Leu Leu Thr Cys Leu Ala  
260 265 270

Leu His Pro Leu Val Val Ile Val Gln Glu Thr Val Glu Ile Thr Glu  
275 280 285

Lys Val Trp Leu Ser Thr Leu Trp Tyr Gln Asn Val Gln Asn  
290 295 300

&lt;210&gt; 143

&lt;211&gt; 1281

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 143

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ttgtttaaga aaaataaaga cattcttaat attgcattgc cagctatggg tgaaaacttt 60
ttgcagatgc taatgggaat ggtggacagt tatttggttg ctcathtagg attgatagct 120
atttcagggg tttcagtagc tggtaatat atcaccattt atcaggcgat tttcatcgct 180
ctgggagctg ctatttccag tgttatttca aaaagcatag ggcagaaaga ccagtcgaag 240
ttggcctatc atgtgactga ggcgttgaag attaccttac tattaagttt ccttttagga 300
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agttttttat ccaatgcctt gaatatctt ttttcaagtc tagctatttt tgttctggat 540
atggggatag ctggtgttg ctgggggaca attgtgtctc gtttggttgg tcttgtgatt 600
ttgtggtcac aattaaaact gccttatggg aagccaactt ttggttttaga taaggaactg 660
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cagggattag gaaatgcacg cctccctttt tatgcgacaa gtataggaat gtggtgtatc 1140
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gcagggtctc tcttgataa tggttttcgc tggttatttc tacgctatcg ttaccagcgc 1260
tatatgagct tgaaaggata g

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&lt;210&gt; 144

&lt;211&gt; 426

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 144

Leu Phe Lys Lys Asn Lys Asp Ile Leu Asn Ile Ala Leu Pro Ala Met

1

5

10

15



Gly Glu Asn Phe Leu Gln Met Leu Met Gly Met Val Asp Ser Tyr Leu  
 20 25 30

Val Ala His Leu Gly Leu Ile Ala Ile Ser Gly Val Ser Val Ala Gly  
 35 40 45

Asn Ile Ile Thr Ile Tyr Gln Ala Ile Phe Ile Ala Leu Gly Ala Ala  
 50 55 60

Ile Ser Ser Val Ile Ser Lys Ser Ile Gly Gln Lys Asp Gln Ser Lys  
 65 70 75 80

Leu Ala Tyr His Val Thr Glu Ala Leu Lys Ile Thr Leu Leu Leu Ser  
 85 90 95

Phe Leu Leu Gly Phe Leu Ser Ile Phe Ala Gly Lys Glu Met Ile Gly  
 100 105 110

Leu Leu Gly Thr Glu Arg Asp Val Ala Glu Ser Gly Gly Leu Tyr Leu  
 115 120 125

Ser Leu Val Gly Gly Ser Ile Val Leu Leu Gly Leu Met Thr Ser Leu  
 130 135 140

Gly Ala Leu Ile Arg Ala Thr His Asn Pro Arg Leu Pro Leu Tyr Val  
 145 150 155 160

Ser Phe Leu Ser Asn Ala Leu Asn Ile Leu Phe Ser Ser Leu Ala Ile  
 165 170 175

Phe Val Leu Asp Met Gly Ile Ala Gly Val Ala Trp Gly Thr Ile Val  
 180 185 190

Ser Arg Leu Val Gly Leu Val Ile Leu Trp Ser Gln Leu Lys Leu Pro  
 195 200 205

Tyr Gly Lys Pro Thr Phe Gly Leu Asp Lys Glu Leu Leu Thr Leu Ala  
 210 215 220

197

Leu Pro Ala Ala Gly Glu Arg Leu Met Met Arg Ala Gly Asp Val Val  
225 230 235 240

Ile Ile Ala Leu Val Val Ser Phe Gly Thr Glu Ala Val Ala Gly Asn  
245 250 255

Ala Ile Gly Glu Val Leu Thr Gln Phe Asn Tyr Met Pro Ala Phe Gly  
260 265 270

Val Ala Thr Ala Thr Val Met Leu Leu Ala Arg Ala Val Gly Glu Asp  
275 280 285

Asp Trp Lys Arg Val Ala Ser Leu Ser Lys Gln Thr Phe Trp Leu Ser  
290 295 300

Leu Phe Leu Met Leu Pro Leu Ser Phe Ser Ile Tyr Val Leu Gly Val  
305 310 315 320

Pro Leu Thr His Leu Tyr Thr Thr Asp Ser Leu Ala Val Glu Ala Ser  
325 330 335

Val Leu Val Thr Leu Phe Ser Leu Leu Gly Thr Pro Met Thr Thr Gly  
340 345 350

Thr Val Ile Tyr Thr Ala Val Trp Gln Gly Leu Gly Asn Ala Arg Leu  
355 360 365

Pro Phe Tyr Ala Thr Ser Ile Gly Met Trp Cys Ile Arg Ile Gly Thr  
370 375 380

Gly Tyr Leu Met Gly Ile Val Leu Gly Trp Gly Leu Pro Gly Ile Trp  
385 390 395 400

Ala Gly Ser Leu Leu Asp Asn Gly Phe Arg Trp Leu Phe Leu Arg Tyr  
405 410 415

Arg Tyr Gln Arg Tyr Met Ser Leu Lys Gly  
420 425

&lt;210&gt; 145

&lt;211&gt; 894

&lt;212&gt; DNA

<213> *Streptococcus pneumoniae*

&lt;400&gt; 145

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tttcgaacaa gtggaaatta ttcttttaaag gaagcagcag gcgaatcctg ctctacctct 120
cagttatctc gctttgagct tggggagtct gacctggcag tctcccgttt ctttgagatt 180
ttggataaca ttcattgtaac aatcgaaaat ttcattggata aggcaaggaa ttttcataat 240
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gatatgaagc aggatgattt gggtaaggta gcagattatc tcttcaaaac agaagaatgg 480
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ttttataatg ccaactattt tgaggcttat acagagaaga ttattgacaa aggtattaag 720
ctttatgagc gtaatgtttt ccattattta aaagggtttg ctttatatca aaaaggacag 780
tgtaaagaag gctgtaagca gatgcaagag gccatgcata tttttgatgt gttaggtctt 840
ccagagcaag tagcctatta tcaggaacac tacgaaaaat ttgtcaaaaag ttaa      894

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&lt;210&gt; 146

&lt;211&gt; 297

&lt;212&gt; PRT

<213> *Streptococcus pneumoniae*

&lt;400&gt; 146

```

Val Gly Arg Ile Ile Arg Ala Gly Val Lys Met Glu His Leu Gly Lys
  1             5             10             15

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Val Phe Arg Glu Phe Arg Thr Ser Gly Asn Tyr Ser Leu Lys Glu Ala
      20             25             30

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Ala Gly Glu Ser Cys Ser Thr Ser Gln Leu Ser Arg Phe Glu Leu Gly  
 35 40 45

Glu Ser Asp Leu Ala Val Ser Arg Phe Phe Glu Ile Leu Asp Asn Ile  
 50 55 60

His Val Thr Ile Glu Asn Phe Met Asp Lys Ala Arg Asn Phe His Asn  
 65 70 75 80

His Glu His Val Ser Met Met Ala Gln Ile Ile Pro Leu Tyr Tyr Ser  
 85 90 95

Asn Asp Ile Ala Gly Phe Gln Lys Leu Gln Arg Glu Gln Leu Glu Lys  
 100 105 110

Ser Lys Ser Ser Thr Thr Pro Leu Tyr Phe Glu Leu Asn Trp Ile Leu  
 115 120 125

Leu Gln Gly Leu Ile Cys Gln Arg Asp Ala Ser Tyr Asp Met Lys Gln  
 130 135 140

Asp Asp Leu Gly Lys Val Ala Asp Tyr Leu Phe Lys Thr Glu Glu Trp  
 145 150 155 160

Thr Met Tyr Glu Leu Ile Leu Phe Gly Asn Leu Tyr Ser Phe Tyr Asp  
 165 170 175

Val Asp Tyr Val Thr Arg Ile Gly Arg Glu Val Met Glu Arg Glu Glu  
 180 185 190

Phe Tyr Gln Glu Ile Ser Arg His Lys Arg Leu Val Leu Ile Leu Ala  
 195 200 205

Leu Asn Cys Tyr Gln His Cys Leu Glu His Ser Ser Phe Tyr Asn Ala  
 210 215 220

Asn Tyr Phe Glu Ala Tyr Thr Glu Lys Ile Ile Asp Lys Gly Ile Lys  
 225 230 235 240

200

Leu Tyr Glu Arg Asn Val Phe His Tyr Leu Lys Gly Phe Ala Leu Tyr  
245 250 255

Gln Lys Gly Gln Cys Lys Glu Gly Cys Lys Gln Met Gln Glu Ala Met  
260 265 270

His Ile Phe Asp Val Leu Gly Leu Pro Glu Gln Val Ala Tyr Tyr Gln  
275 280 285

Glu His Tyr Glu Lys Phe Val Lys Ser  
290 295

<210> 147

<211> 1068

<212> DNA

<213> Streptococcus pneumoniae

<400> 147

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cagcgccgta ttctttattc tatgaataag gatagcaata cttttgacaa gagctaccgt 180  
aagtcggcca agtcagtcgg gaacatcatg gggaatttcc acccacacgg ggattcttct 240  
atctatgatg ccatggttcg tatgtcacag aactggaaaa atcgtgagat tctagttgaa 300  
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gcacgtttgt ctgaaattgc aggctacctt cttcaggata tcgagaaaaa gacagttcct 420  
tttgcatgga actttgacga tacggagaaa gaaccaacgg tcttgccagc agcctttcca 480  
aacctcttgg tcaatgggtc gactgggatt tcggctgggt atgccacaga cattcctccc 540  
cataatttag ctgaggtcat agatgctgca gtttacatga ttgaccaccc aactgcaaag 600  
attgataaac tcatggaatt cttgcctgga ccagacttcc ctacaggggc tattattcag 660  
ggtcgtgatg aaatcaagaa agcttatgag actgggaaaag ggcgcggtgt tgttcgttcc 720  
aagactgaaa ttgaaaagct aaaaggtggt aaggaacaaa tcgttattat tgagattcct 780  
tatgaaatca ataaggccaa tctagtcaag aaaatcgatg atgttcgtgt taataacaag 840  
gtagctggga ttgctgaggt tcgtgatgag tctgaccgtg atggtcttcg tctcgtatc 900  
gaacttaaga aagacgctaa tactgagctt gttctcaact acttatttaa gtacaccgac 960  
ctacaaatca actacaactt taatatggtg gcgattgaca atttcacacc tcgtcagggt 1020  
ggattgttcc aatcctgtct agctatatcg ctcaccgtcg agaagtga 1068

&lt;210&gt; 148

&lt;211&gt; 355

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 148

Met Ser Asn Ile Gln Asn Met Ser Leu Glu Asp Ile Met Gly Glu Arg

1

5

10

15

Phe Gly Arg Tyr Ser Lys Tyr Ile Ile Gln Asp Arg Ala Leu Pro Asp

20

25

30

Ile Arg Asp Gly Leu Lys Pro Val Gln Arg Arg Ile Leu Tyr Ser Met

35

40

45

Asn Lys Asp Ser Asn Thr Phe Asp Lys Ser Tyr Arg Lys Ser Ala Lys

50

55

60

Ser Val Gly Asn Ile Met Gly Asn Phe His Pro His Gly Asp Ser Ser

65

70

75

80

Ile Tyr Asp Ala Met Val Arg Met Ser Gln Asn Trp Lys Asn Arg Glu

85

90

95

Ile Leu Val Glu Met His Gly Asn Asn Gly Ser Met Asp Gly Asp Pro

100

105

110

Pro Ala Ala Met Arg Tyr Thr Glu Ala Arg Leu Ser Glu Ile Ala Gly

115

120

125

Tyr Leu Leu Gln Asp Ile Glu Lys Lys Thr Val Pro Phe Ala Trp Asn

130

135

140

Phe Asp Asp Thr Glu Lys Glu Pro Thr Val Leu Pro Ala Ala Phe Pro

145

150

155

160

202

Asn Leu Leu Val Asn Gly Ser Thr Gly Ile Ser Ala Gly Tyr Ala Thr  
165 170 175

Asp Ile Pro Pro His Asn Leu Ala Glu Val Ile Asp Ala Ala Val Tyr  
180 185 190

Met Ile Asp His Pro Thr Ala Lys Ile Asp Lys Leu Met Glu Phe Leu  
195 200 205

Pro Gly Pro Asp Phe Pro Thr Gly Ala Ile Ile Gln Gly Arg Asp Glu  
210 215 220

Ile Lys Lys Ala Tyr Glu Thr Gly Lys Gly Arg Val Val Val Arg Ser  
225 230 235 240

Lys Thr Glu Ile Glu Lys Leu Lys Gly Gly Lys Glu Gln Ile Val Ile  
245 250 255

Ile Glu Ile Pro Tyr Glu Ile Asn Lys Ala Asn Leu Val Lys Lys Ile  
260 265 270

Asp Asp Val Arg Val Asn Asn Lys Val Ala Gly Ile Ala Glu Val Arg  
275 280 285

Asp Glu Ser Asp Arg Asp Gly Leu Arg Ile Ala Ile Glu Leu Lys Lys  
290 295 300

Asp Ala Asn Thr Glu Leu Val Leu Asn Tyr Leu Phe Lys Tyr Thr Asp  
305 310 315 320

Leu Gln Ile Asn Tyr Asn Phe Asn Met Val Ala Ile Asp Asn Phe Thr  
325 330 335

Pro Arg Gln Val Gly Leu Phe Gln Ser Cys Leu Ala Ile Ser Leu Thr  
340 345 350

Val Glu Lys  
355

&lt;210&gt; 149

&lt;211&gt; 684

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 149

```

atgccgacat tagaaatagc acaaaaaaaaa ctggagttca ttaagaaggc agaagaatat 60
tacaatgcct tgtgtacaaa tatacagttg agcggagata aactaaaagt aatttccggt 120
acttctgtta accctgggga aggaaaaaca actacttcca taaatatagc atggtcgttt 180
gcgcggtgcag gctataaaac tcttttgatc gatggcgata ctcgaaattc agttatgtta 240
ggagttttta aatctcgtga aaaaattaca gggctaacag aatttttata tgggacagct 300
gatttatctc acggtttatg tgatacaaat attgaaaatt tatttgtagt tcaatcggga 360
tctgtatcac caaacctac agccttgta caaagtaaaa attttaatga tatgattgaa 420
acattgcgta aatattttga ttatatcatt attgatacac cgcctattgg aattgttatt 480
gatgcggcaa ttatcactca aaagtgtgat gcgtccatct tggtaacagc aacaggtgag 540
gcgaataaac gtgatatcca aaaagcgaaa caacaattaa aacaaacagg gaaactgttc 600
ctaggagttg ttttaaataa attggatatc tcggttaata agtatggagt ttacggttcc 660
tatggaaatt atggtaaaaa ataa                                     684

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&lt;210&gt; 150

&lt;211&gt; 227

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 150

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Met Pro Thr Leu Glu Ile Ala Gln Lys Lys Leu Glu Phe Ile Lys Lys
  1             5             10            15

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```

Ala Glu Glu Tyr Tyr Asn Ala Leu Cys Thr Asn Ile Gln Leu Ser Gly
      20             25            30

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Asp Lys Leu Lys Val Ile Ser Val Thr Ser Val Asn Pro Gly Glu Gly
      35             40            45

```



Gly Lys Lys  
225

&lt;210&gt; 151

&lt;211&gt; 1194

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 151

```

atggaggcaa atatgaaaca tctaaaaaca ttttacaaaa aatgggtttca attattagtc 60
gttatcgtca ttagcttttt tagtggagcc ttgggtagtt tttcaataac tcaactaact 120
caaaaaagta gtgtaaacaa ctctaacaac aatagtacta ttacacaaac tgcctataag 180
aacgaaaatt caacaacaca ggctgttaac aaagtaaaag atgctgttgt ttctgttatt 240
acttattcgg caaacagaca aaatagcgta tttggcaatg atgatactga cacagattct 300
cagcgaatct ctagtgaagg atctggagtt atttataaaa agaagcttac 360
atcgtcacca acaatcacgt tattaatggc gccagcaaag tagatattcg attgtcagat 420
gggactaaag tacctggaga aattgtcgga gctgacactt tctctgatat tgctgtcgtc 480
aaaatctctt cagaaaaaagt gacaacagta gctgagtttg gtgattctag taagttaact 540
gtaggagaaa ctgctattgc catcggtagc ccggttaggt ctgaatatgc aaatactgtc 600
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gctattttcta caaaagccat ccaaactgat actgctatta acccaggtaa ctctggcggc 720
ccactgatca atattcaagg gcagggtatc ggaattacct caagtaaaat tgctacaaat 780
ggaggaacat ctgtagaagg tcttggtttc gcaattcctg caaatgatgc tatcaatatt 840
attgaacagt tagaaaaaaa cggaaaagtg acgcgtccag ctttgggaat ccagatgggt 900
aatttatcta atgtgagtac aagcgacatc agaagactca atattccaag taatgttaca 960
totggtgtaa ttgttcgttc ggtacaaagt aatatgctg ccaatggtca ccttgaaaaa 1020
tacgatgtaa ttacaaaagt agatgacaaa gagattgott catcaacaga cttacaaagt 1080
gctctttaca accattctat cggagacacc attaagataa cctactatcg taacgggaaa 1140
gaagaaaacta cctctatcaa acttaacaag agttcaggtg atttagaatc ttaa      1194

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&lt;210&gt; 152

&lt;211&gt; 397

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 152

Met Glu Ala Asn Met Lys His Leu Lys Thr Phe Tyr Lys Lys Trp Phe

1

5

10

15

Gln Leu Leu Val Val Ile Val Ile Ser Phe Phe Ser Gly Ala Leu Gly  
 20 25 30

Ser Phe Ser Ile Thr Gln Leu Thr Gln Lys Ser Ser Val Asn Asn Ser  
 35 40 45

Asn Asn Asn Ser Thr Ile Thr Gln Thr Ala Tyr Lys Asn Glu Asn Ser  
 50 55 60

Thr Thr Gln Ala Val Asn Lys Val Lys Asp Ala Val Val Ser Val Ile  
 65 70 75 80

Thr Tyr Ser Ala Asn Arg Gln Asn Ser Val Phe Gly Asn Asp Asp Thr  
 85 90 95

Asp Thr Asp Ser Gln Arg Ile Ser Ser Glu Gly Ser Gly Val Ile Tyr  
 100 105 110

Lys Lys Asn Asp Lys Glu Ala Tyr Ile Val Thr Asn Asn His Val Ile  
 115 120 125

Asn Gly Ala Ser Lys Val Asp Ile Arg Leu Ser Asp Gly Thr Lys Val  
 130 135 140

Pro Gly Glu Ile Val Gly Ala Asp Thr Phe Ser Asp Ile Ala Val Val  
 145 150 155 160

Lys Ile Ser Ser Glu Lys Val Thr Thr Val Ala Glu Phe Gly Asp Ser  
 165 170 175

Ser Lys Leu Thr Val Gly Glu Thr Ala Ile Ala Ile Gly Ser Pro Leu  
 180 185 190

Gly Ser Glu Tyr Ala Asn Thr Val Thr Gln Gly Ile Val Ser Ser Leu  
 195 200 205

Asn Arg Asn Val Ser Leu Lys Ser Glu Asp Gly Gln Ala Ile Ser Thr  
 210 215 220

207

Lys Ala Ile Gln Thr Asp Thr Ala Ile Asn Pro Gly Asn Ser Gly Gly  
225 230 235 240

Pro Leu Ile Asn Ile Gln Gly Gln Val Ile Gly Ile Thr Ser Ser Lys  
245 250 255

Ile Ala Thr Asn Gly Gly Thr Ser Val Glu Gly Leu Gly Phe Ala Ile  
260 265 270

Pro Ala Asn Asp Ala Ile Asn Ile Ile Glu Gln Leu Glu Lys Asn Gly  
275 280 285

Lys Val Thr Arg Pro Ala Leu Gly Ile Gln Met Val Asn Leu Ser Asn  
290 295 300

Val Ser Thr Ser Asp Ile Arg Arg Leu Asn Ile Pro Ser Asn Val Thr  
305 310 315 320

Ser Gly Val Ile Val Arg Ser Val Gln Ser Asn Met Pro Ala Asn Gly  
325 330 335

His Leu Glu Lys Tyr Asp Val Ile Thr Lys Val Asp Asp Lys Glu Ile  
340 345 350

Ala Ser Ser Thr Asp Leu Gln Ser Ala Leu Tyr Asn His Ser Ile Gly  
355 360 365

Asp Thr Ile Lys Ile Thr Tyr Tyr Arg Asn Gly Lys Glu Glu Thr Thr  
370 375 380

Ser Ile Lys Leu Asn Lys Ser Ser Gly Asp Leu Glu Ser  
385 390 395

&lt;210&gt; 153

&lt;211&gt; 939

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 153

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atggcagaaa tttatctagc aggtggttgt ttttggggcc tagaggaata tttttcacgc 60
atttctggag tgctagaaac cagtgttggc tacgctaata gtcaagtcga aacgaccaat 120
taccagttgc tcaaggaaac agaccatgca gaaacggtcc aagtgattta cgatgagaag 180
gaagtgtcac tcagagagat tttactttat tttttccgag ttatcgatcc tctatctatc 240
aatcaacaag ggaatgaccg tggtcgccaa tatcgaaact ggatttatta tcaggatgaa 300
gcagatttgc cagctatcta cacagtgggt caggagcagg aacgcatgct gggtcgaaaag 360
attgcagtag aagtggagca attacgccac tacattctgg ctgaagacta ccaccaagac 420
tatctcagga agaatccttc aggttactgt catatcgatg tgaccgatgc tgataagcca 480
ttgattgatg cagcaaaacta tgaaaagcct agtcaagagg tgttgaaggc cagtctatct 540
gaagagtctt atcgtgtcac acaagaagct gctacagagg ctccatttac caatgcctat 600
gaccaaactt ttgaagaggg gatttatgta gatattacga caggtgagcc actctttttt 660
gccaaggata agtttgcttc aggttgttgt tggccaagtt ttagccgtcc gatttccaaa 720
gagttgattc attattacaa ggatctgagc catggaatgg agcgaattga agttcgttct 780
cgttcaggca gtgctcactt gggtcatgtt ttcacagatg gaccgcggga gttaggcggc 840
ctccgttact gtatcaattc tgcttcttta cgctttgtgg ccaaggatga gatggaaaaa 900
gcaggatatg gctatctatt gccttactta aacaaaataa 939

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&lt;210&gt; 154

&lt;211&gt; 312

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 154

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Met Ala Glu Ile Tyr Leu Ala Gly Gly Cys Phe Trp Gly Leu Glu Glu
  1             5             10             15

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```

Tyr Phe Ser Arg Ile Ser Gly Val Leu Glu Thr Ser Val Gly Tyr Ala
      20             25             30

```

```

Asn Gly Gln Val Glu Thr Thr Asn Tyr Gln Leu Leu Lys Glu Thr Asp
      35             40             45

```

His Ala Glu Thr Val Gln Val Ile Tyr Asp Glu Lys Glu Val Ser Leu  
 50 55 60

Arg Glu Ile Leu Leu Tyr Tyr Phe Arg Val Ile Asp Pro Leu Ser Ile  
 65 70 75 80

Asn Gln Gln Gly Asn Asp Arg Gly Arg Gln Tyr Arg Thr Gly Ile Tyr  
 85 90 95

Tyr Gln Asp Glu Ala Asp Leu Pro Ala Ile Tyr Thr Val Val Gln Glu  
 100 105 110

Gln Glu Arg Met Leu Gly Arg Lys Ile Ala Val Glu Val Glu Gln Leu  
 115 120 125

Arg His Tyr Ile Leu Ala Glu Asp Tyr His Gln Asp Tyr Leu Arg Lys  
 130 135 140

Asn Pro Ser Gly Tyr Cys His Ile Asp Val Thr Asp Ala Asp Lys Pro  
 145 150 155 160

Leu Ile Asp Ala Ala Asn Tyr Glu Lys Pro Ser Gln Glu Val Leu Lys  
 165 170 175

Ala Ser Leu Ser Glu Glu Ser Tyr Arg Val Thr Gln Glu Ala Ala Thr  
 180 185 190

Glu Ala Pro Phe Thr Asn Ala Tyr Asp Gln Thr Phe Glu Glu Gly Ile  
 195 200 205

Tyr Val Asp Ile Thr Thr Gly Glu Pro Leu Phe Phe Ala Lys Asp Lys  
 210 215 220

Phe Ala Ser Gly Cys Gly Trp Pro Ser Phe Ser Arg Pro Ile Ser Lys  
 225 230 235 240

Glu Leu Ile His Tyr Tyr Lys Asp Leu Ser His Gly Met Glu Arg Ile  
 245 250 255

210

Glu Val Arg Ser Arg Ser Gly Ser Ala His Leu Gly His Val Phe Thr  
260 265 270

Asp Gly Pro Arg Glu Leu Gly Gly Leu Arg Tyr Cys Ile Asn Ser Ala  
275 280 285

Ser Leu Arg Phe Val Ala Lys Asp Glu Met Glu Lys Ala Gly Tyr Gly  
290 295 300

Tyr Leu Leu Pro Tyr Leu Asn Lys  
305 310

<210> 155

<211> 870

<212> DNA

<213> Streptococcus pneumoniae

<400> 155

atgaagatta ttgtacctgc aaccagtgcc aatatcgggc caggttttga ctcggtcgggt 60  
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gaacaccaga ttggcaaattg gattccacat gacgagcgta atctcttgct caaaatcgct 180  
ttgcaaattg taccagactt gcaaccaaga cgcttgaaaa tgaccagtga tgtccctttg 240  
gcgcgcggtt tgggttcttc cagctcgggt atcgttgctg ggattgaact agccaaccaa 300  
ctgggtcaac tcaacttatc agaccatgaa aaattgcagt tagcgaccaa gattgaaggg 360  
catcctgaca atgtggctcc agccatttat ggtaatctcg ttattgcaag ttctgttgaa 420  
gggcaagtct ctgctatcgt agcagacttt ccagagtgtg attttctagc ttacattcca 480  
aactatgaat tacgtactcg cgacagccgt agtgtcttgc ctaaaaaatt gtcttataag 540  
gaagctgttg ctgcaagttc tatcgccaat gtagcgggtg ctgccttggt ggcaggagac 600  
atggtgaccg ctgggcaagc aatcgaggga gacctcttcc atgagcgcta tcgtcaggac 660  
ttggtaagag aatttgcatg gattaagcaa gtgaccaaag aaaatggggc ctatgcaacc 720  
tacctttctg gtgctggggc gacagttatg gttctggctt ctcatgacaa gatgccaaca 780  
attaaggcag aattggaaaa gcaacctttc aaaggaaaac tgcattgactt gagagttgat 840  
acccaagggtg tccgtgtaga agcaaaaataa 870

&lt;210&gt; 156

&lt;211&gt; 289

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 156

Met Lys Ile Ile Val Pro Ala Thr Ser Ala Asn Ile Gly Pro Gly Phe

1

5

10

15

Asp Ser Val Gly Val Ala Val Thr Lys Tyr Leu Gln Ile Glu Val Cys

20

25

30

Glu Glu Arg Asp Glu Trp Leu Ile Glu His Gln Ile Gly Lys Trp Ile

35

40

45

Pro His Asp Glu Arg Asn Leu Leu Leu Lys Ile Ala Leu Gln Ile Val

50

55

60

Pro Asp Leu Gln Pro Arg Arg Leu Lys Met Thr Ser Asp Val Pro Leu

65

70

75

80

Ala Arg Gly Leu Gly Ser Ser Ser Ser Val Ile Val Ala Gly Ile Glu

85

90

95

Leu Ala Asn Gln Leu Gly Gln Leu Asn Leu Ser Asp His Glu Lys Leu

100

105

110

Gln Leu Ala Thr Lys Ile Glu Gly His Pro Asp Asn Val Ala Pro Ala

115

120

125

Ile Tyr Gly Asn Leu Val Ile Ala Ser Ser Val Glu Gly Gln Val Ser

130

135

140

Ala Ile Val Ala Asp Phe Pro Glu Cys Asp Phe Leu Ala Tyr Ile Pro

145

150

155

160

Asn Tyr Glu Leu Arg Thr Arg Asp Ser Arg Ser Val Leu Pro Lys Lys

165

170

175



212

Leu Ser Tyr Lys Glu Ala Val Ala Ala Ser Ser Ile Ala Asn Val Ala  
180 185 190

Val Ala Ala Leu Leu Ala Gly Asp Met Val Thr Ala Gly Gln Ala Ile  
195 200 205

Glu Gly Asp Leu Phe His Glu Arg Tyr Arg Gln Asp Leu Val Arg Glu  
210 215 220

Phe Ala Met Ile Lys Gln Val Thr Lys Glu Asn Gly Ala Tyr Ala Thr  
225 230 235 240

Tyr Leu Ser Gly Ala Gly Pro Thr Val Met Val Leu Ala Ser His Asp  
245 250 255

Lys Met Pro Thr Ile Lys Ala Glu Leu Glu Lys Gln Pro Phe Lys Gly  
260 265 270

Lys Leu His Asp Leu Arg Val Asp Thr Gln Gly Val Arg Val Glu Ala  
275 280 285

Lys

<210> 157

<211> 564

<212> DNA

<213> Streptococcus pneumoniae

<400> 157

atgaaatatac acgattacat ctgggattta ggtggaaactt tactggataa ttatgaaact 60  
tcaacagctg catttgttga aacattggca ctgtatggta tcacacaaga ccatgacagt 120  
gtctatcaag ctttaaaggt ttctactcct tttgcgattg agacattcgc tcccaattta 180  
gagaatTTTT tagaaaagta caaggaaaat gaagccagag agcttgaaca cccgatttta 240  
tttgaaggag tttctgacct attggaagac atttcaaatc aaggtggccg tcattttttg 300  
gtctctcatc gaaatgatca ggTTTTggaa atttttagaaa aaacctotat agcagcttat 360

tttacagaag tgggtgacttc tagctcaggc tttaagagaa agccaaatcc cgaatccatg 420  
 ctttatattaa gagaaaagta tcagattagc tctggtcttg tcattggtga toggcgcgatt 480  
 gatatacgaag caggtcaagc tgcaggactt gataccact tgtttaccag tatcgtgaat 540  
 ttaagacaag tattagacat ataa 564

<210> 158

<211> 187

<212> PRT

<213> Streptococcus pneumoniae

<400> 158

Met Lys Tyr His Asp Tyr Ile Trp Asp Leu Gly Gly Thr Leu Leu Asp  
 1 5 10 15

Asn Tyr Glu Thr Ser Thr Ala Ala Phe Val Glu Thr Leu Ala Leu Tyr  
 20 25 30

Gly Ile Thr Gln Asp His Asp Ser Val Tyr Gln Ala Leu Lys Val Ser  
 35 40 45

Thr Pro Phe Ala Ile Glu Thr Phe Ala Pro Asn Leu Glu Asn Phe Leu  
 50 55 60

Glu Lys Tyr Lys Glu Asn Glu Ala Arg Glu Leu Glu His Pro Ile Leu  
 65 70 75 80

Phe Glu Gly Val Ser Asp Leu Leu Glu Asp Ile Ser Asn Gln Gly Gly  
 85 90 95

Arg His Phe Leu Val Ser His Arg Asn Asp Gln Val Leu Glu Ile Leu  
 100 105 110

Glu Lys Thr Ser Ile Ala Ala Tyr Phe Thr Glu Val Val Thr Ser Ser  
 115 120 125

Ser Gly Phe Lys Arg Lys Pro Asn Pro Glu Ser Met Leu Tyr Leu Arg  
 130 135 140

Glu Lys Tyr Gln Ile Ser Ser Gly Leu Val Ile Gly Asp Arg Pro Ile  
 145 150 155 160

Asp Ile Glu Ala Gly Gln Ala Ala Gly Leu Asp Thr His Leu Phe Thr  
 165 170 175

Ser Ile Val Asn Leu Arg Gln Val Leu Asp Ile  
 180 185

<210> 159

<211> 1875

<212> DNA

<213> Streptococcus pneumoniae

<400> 159

atgacagaag aaatcaaaaa tctgcaggca caggattatg atgccagtca aattcaagtt 60  
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 gaaggtcttc accatctagt ctgggaaatt gttgataact caattgacga ggccttggca 180  
 ggatttgcca gccatattca agtttttatt gagccagatg attogattac tgttgtggat 240  
 gatgggcgtg gtatcccagt cgatattcag gaaaaaacag gccgtcctgc tgttgagacc 300  
 gtctttacag tccttcacgc tggaggaaaag ttcggcgggtg gtggatacaa ggtttcaggt 360  
 ggtcttcacg ggggtggggtc gtcagtagtt aatgcccttt ccaactcaatt agacgttcat 420  
 gttcacaaaa atggtaagat tcattaccaa gaataccgtc gtgggtcatgt tgtcgcagat 480  
 cttgaaatag ttggagatac ggataaaaca ggaacaactg ttcaacttcac accggaccca 540  
 aaaatcttca ctgaaacaac aatctttgat tttgataaat taaataaacg gattcaagag 600  
 ttggcctttc taaatcgcgg tcttcaaatt tcaattacag ataagcgcca aggtttggaa 660  
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 aacaaggatg taatctttga tacaccaatc tatacagacg gtgagatgga tgatatcaca 780  
 gttgaggtag ccatgcagta cacaactggg taccatgaaa atgtcatgag tttcgccaat 840  
 aatattcata cccatgaagg tggaacacat gaacaagggt tccgtacagc cttgacacgt 900  
 gttatcaacg attatgctcg taaaaataag ttactgaaag acaatgaaga taatttaaca 960  
 ggggaagatg ttcgcgaagg cttaactgca gttatctcag ttaaacaccc aaatccacag 1020  
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 gtagaaaaag gaattttggc tgccaagggt cgtgtgggtg ccaagcgtgc gcgtgaagtc 1200  
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 gccaaatctg gtcgtaaccg tgagtttcag gctatccttc caattcgcgg taagattttg 1380  
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 acagccatgg gaacaggatt tggcgcagaa tttgatgttt cgaaagcccg ttacccaaaaa 1500  
 ctcgttttga tgaccgatgc cgatgtcgat ggagcccaca ttcgtaccct tottttaacc 1560  
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 ccaatctatg gtgtcaaggt tggaagcgag attaaagaat atatccagcc ggggtgcagat 1680  
 caagaaatca aactccaaga agctttagcc cgttatagtg aaggctgtac caaaccgact 1740  
 attcagcgtt ataaggggct aggtgaaatg gacgatcatc agctgtggga aacaaccatg 1800  
 gatcccgaac atcgcttgat ggctagagtt tctgtagatg atgtgcagaa gcagataaaaa 1860  
 tctttgatat gttga 1875

<210> 160

<211> 624

<212> PRT

<213> Streptococcus pneumoniae

<400> 160

Met Thr Glu Glu Ile Lys Asn Leu Gln Ala Gln Asp Tyr Asp Ala Ser  
 1 5 10 15

Gln Ile Gln Val Leu Glu Gly Leu Glu Ala Val Arg Met Arg Pro Gly  
 20 25 30

Met Tyr Ile Gly Ser Thr Ser Lys Glu Gly Leu His His Leu Val Trp  
 35 40 45

Glu Ile Val Asp Asn Ser Ile Asp Glu Ala Leu Ala Gly Phe Ala Ser  
 50 55 60

His Ile Gln Val Phe Ile Glu Pro Asp Asp Ser Ile Thr Val Val Asp  
 65 70 75 80

Asp Gly Arg Gly Ile Pro Val Asp Ile Gln Glu Lys Thr Gly Arg Pro  
 85 90 95

216

Ala Val Glu Thr Val Phe Thr Val Leu His Ala Gly Gly Lys Phe Gly  
100 105 110

Gly Gly Gly Tyr Lys Val Ser Gly Gly Leu His Gly Val Gly Ser Ser  
115 120 125

Val Val Asn Ala Leu Ser Thr Gln Leu Asp Val His Val His Lys Asn  
130 135 140

Gly Lys Ile His Tyr Gln Glu Tyr Arg Arg Gly His Val Val Ala Asp  
145 150 155 160

Leu Glu Ile Val Gly Asp Thr Asp Lys Thr Gly Thr Thr Val His Phe  
165 170 175

Thr Pro Asp Pro Lys Ile Phe Thr Glu Thr Thr Ile Phe Asp Phe Asp  
180 185 190

Lys Leu Asn Lys Arg Ile Gln Glu Leu Ala Phe Leu Asn Arg Gly Leu  
195 200 205

Gln Ile Ser Ile Thr Asp Lys Arg Gln Gly Leu Glu Gln Thr Lys His  
210 215 220

Tyr His Tyr Glu Gly Gly Ile Ala Ser Tyr Val Glu Tyr Ile Asn Glu  
225 230 235 240

Asn Lys Asp Val Ile Phe Asp Thr Pro Ile Tyr Thr Asp Gly Glu Met  
245 250 255

Asp Asp Ile Thr Val Glu Val Ala Met Gln Tyr Thr Thr Gly Tyr His  
260 265 270

Glu Asn Val Met Ser Phe Ala Asn Asn Ile His Thr His Glu Gly Gly  
275 280 285

Thr His Glu Gln Gly Phe Arg Thr Ala Leu Thr Arg Val Ile Asn Asp  
290 295 300

Variable	Unit	Mean	SD	Min	Max
Age	Years	45.2	12.5	25	65
Gender	Male/Female	50/50			
Education	Years	12.8	2.1	8	16
Income	\$/month	1500	500	500	3000
Marital status	Married/Single	60/40			
Occupation	Various				
Health status	Good/Poor	70/30			
Smoking status	Smoker/Non-smoker	30/70			
Alcohol consumption	Yes/No	20/80			
Family size	Number	3.5	1.2	1	6
Urban/rural	Urban/Rural	65/35			
Religion	Various				
Political affiliation	Various				
Language spoken	Various				
Health insurance	Yes/No	80/20			
Access to healthcare	Yes/No	90/10			
Healthcare costs	\$/year	200	100	0	500
Healthcare quality	Good/Poor	75/25			
Healthcare access	Yes/No	85/15			
Healthcare satisfaction	Yes/No	70/30			
Healthcare utilization	Yes/No	80/20			
Healthcare expenditure	\$/year	150	75	0	300
Healthcare coverage	Yes/No	95/5			
Healthcare quality of care	Good/Poor	80/20			
Healthcare access to services	Yes/No	90/10			
Healthcare satisfaction with care	Yes/No	75/25			
Healthcare utilization of services	Yes/No	85/15			
Healthcare expenditure on services	\$/year	100	50	0	200
Healthcare coverage of services	Yes/No	90/10			
Healthcare quality of services	Good/Poor	85/15			
Healthcare access to specialists	Yes/No	80/20			
Healthcare satisfaction with specialists	Yes/No	70/30			
Healthcare utilization of specialists	Yes/No	75/25			
Healthcare expenditure on specialists	\$/year	50	25	0	100
Healthcare coverage of specialists	Yes/No	85/15			
Healthcare quality of specialists	Good/Poor	90/10			
Healthcare access to primary care	Yes/No	95/5			
Healthcare satisfaction with primary care	Yes/No	80/20			
Healthcare utilization of primary care	Yes/No	90/10			
Healthcare expenditure on primary care	\$/year	50	25	0	100
Healthcare coverage of primary care	Yes/No	95/5			
Healthcare quality of primary care	Good/Poor	90/10			

Arg Tyr Gln Lys Leu Val Leu Met Thr Asp Ala Asp Val Asp Gly Ala  
500 505 510

His Ile Arg Thr Leu Leu Leu Thr Leu Ile Tyr Arg Tyr Met Lys Pro  
 515 520 525

Ile Leu Glu Ala Gly Tyr Val Tyr Ile Ala Gln Pro Pro Ile Tyr Gly  
 530 535 540

Val Lys Val Gly Ser Glu Ile Lys Glu Tyr Ile Gln Pro Gly Ala Asp  
 545 550 555 560

Gln Glu Ile Lys Leu Gln Glu Ala Leu Ala Arg Tyr Ser Glu Gly Arg  
 565 570 575

Thr Lys Pro Thr Ile Gln Arg Tyr Lys Gly Leu Gly Glu Met Asp Asp  
 580 585 590

His Gln Leu Trp Glu Thr Thr Met Asp Pro Glu His Arg Leu Met Ala  
 595 600 605

Arg Val Ser Val Asp Asp Val Gln Lys Gln Ile Lys Ser Leu Ile Cys  
 610 615 620

<210> 161

<211> 1446

<212> DNA

<213> Streptococcus pneumoniae

<400> 161

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&lt;210&gt; 162

&lt;211&gt; 481

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 162

Met Ser Arg Arg Phe Lys Lys Ser Arg Ser Gln Lys Val Lys Arg Ser  
1 5 10 15

Val Asn Ile Val Leu Leu Thr Ile Tyr Leu Leu Leu Val Cys Phe Leu  
20 25 30

Leu Phe Leu Ile Phe Lys Tyr Asn Ile Leu Ala Phe Arg Tyr Leu Asn  
35 40 45

Leu Val Val Thr Ala Leu Val Leu Leu Val Ala Leu Val Gly Leu Leu  
50 55 60



220

Leu Ile Ile Tyr Lys Lys Ala Glu Lys Phe Thr Ile Phe Leu Leu Val  
65 70 75 80

Phe Ser Ile Leu Val Ser Ser Val Ser Leu Phe Ala Val Gln Gln Phe  
85 90 95

Val Gly Leu Thr Asn Arg Leu Asn Ala Thr Ser Asn Tyr Ser Glu Tyr  
100 105 110

Ser Ile Ser Val Ala Val Leu Ala Asp Ser Glu Ile Glu Asn Val Thr  
115 120 125

Gln Leu Thr Ser Val Thr Ala Pro Thr Gly Thr Asn Asn Glu Asn Ile  
130 135 140

Gln Lys Leu Leu Ala Asp Ile Lys Ser Ser Gln Asn Thr Asp Leu Thr  
145 150 155 160

Val Asn Gln Ser Ser Ser Tyr Leu Ala Ala Tyr Lys Ser Leu Ile Ala  
165 170 175

Gly Glu Thr Lys Ala Ile Val Leu Asn Ser Val Phe Glu Asn Ile Ile  
180 185 190

Glu Ser Glu Tyr Pro Asp Tyr Ala Ser Lys Ile Lys Lys Ile Tyr Thr  
195 200 205

Lys Gly Phe Thr Lys Lys Val Glu Ala Pro Lys Thr Ser Lys Ser Gln  
210 215 220

Ser Phe Asn Ile Tyr Val Ser Gly Ile Asp Thr Tyr Gly Pro Ile Ser  
225 230 235 240

Ser Val Ser Arg Ser Asp Val Asn Ile Leu Met Thr Val Asn Arg Asp  
245 250 255

Thr Lys Lys Ile Leu Leu Thr Thr Thr Pro Arg Asp Ala Tyr Val Pro  
260 265 270

Ile Ala Asp Gly Gly Asn Asn Gln Lys Asp Lys Leu Thr His Ala Gly  
 275 280 285

Ile Tyr Gly Val Asp Ser Ser Ile His Thr Leu Glu Asn Leu Tyr Gly  
 290 295 300

Val Asp Ile Asn Tyr Tyr Val Arg Leu Asn Phe Thr Ser Phe Leu Lys  
 305 310 315 320

Leu Ile Asp Leu Leu Gly Gly Ile Asp Val Tyr Asn Asp Gln Glu Phe  
 325 330 335

Thr Ala His Thr Asn Gly Lys Tyr Tyr Pro Ala Gly Asn Val His Leu  
 340 345 350

Asp Ser Glu Gln Ala Leu Gly Phe Val Arg Glu Arg Tyr Ser Leu Ala  
 355 360 365

Asp Gly Asp Arg Asp Arg Gly Arg His Gln Gln Lys Val Ile Val Ala  
 370 375 380

Ile Leu Gln Lys Leu Thr Ser Thr Glu Val Leu Lys Asn Tyr Ser Thr  
 385 390 395 400

Ile Ile Asn Ser Leu Gln Asp Ser Ile Gln Thr Asn Met Pro Leu Glu  
 405 410 415

Thr Met Ile Asn Leu Val Asn Ala Gln Leu Glu Ser Gly Gly Asn Tyr  
 420 425 430

Lys Val Asn Ser Gln Asp Leu Lys Gly Thr Gly Arg Met Asp Leu Pro  
 435 440 445

Ser Tyr Ala Met Pro Asp Ser Asn Leu Tyr Val Met Glu Ile Asp Asp  
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Ser Ser Leu Ala Val Val Lys Ala Ala Ile Gln Asp Val Met Glu Gly  
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Arg

&lt;210&gt; 163

&lt;211&gt; 732

&lt;212&gt; DNA

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 163

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&lt;210&gt; 164

&lt;211&gt; 243

&lt;212&gt; PRT

&lt;213&gt; Streptococcus pneumoniae

&lt;400&gt; 164

Met Ile Asp Ile His Ser His Ile Val Phe Asp Val Asp Asp Gly Pro  
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Lys Ser Arg Glu Glu Ser Lys Ala Leu Leu Ala Glu Ser Tyr Arg Gln  
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Gly Val Arg Thr Ile Val Ser Thr Ser His Arg Arg Lys Gly Met Phe  
 35 40 45

Glu Thr Pro Glu Glu Lys Ile Ala Glu Asn Phe Leu Gln Val Arg Glu  
 50 55 60

Ile Ala Lys Glu Val Ala Ser Asp Leu Val Ile Ala Tyr Gly Ala Glu  
 65 70 75 80

Ile Tyr Tyr Thr Pro Asp Val Leu Asp Lys Leu Glu Lys Lys Arg Ile  
 85 90 95

Pro Thr Leu Asn Asp Ser Arg Tyr Ala Leu Ile Glu Phe Ser Met Asn  
 100 105 110

Thr Pro Tyr Arg Asp Ile His Ser Ala Leu Ser Lys Ile Leu Met Leu  
 115 120 125

Gly Ile Thr Pro Val Ile Ala His Ile Glu Arg Tyr Asp Ala Leu Glu  
 130 135 140

Asn Asn Glu Lys Arg Val Arg Glu Leu Ile Asp Met Gly Cys Tyr Thr  
 145 150 155 160

Gln Val Asn Ser Ser His Val Leu Lys Pro Lys Leu Phe Gly Glu Arg  
 165 170 175

Tyr Lys Phe Met Lys Lys Arg Ala Gln Tyr Phe Leu Glu Gln Asp Leu  
 180 185 190

Val His Val Ile Ala Ser Asp Met His Asn Leu Asp Gly Arg Pro Pro  
 195 200 205

His Met Ala Glu Ala Tyr Asp Leu Val Thr Gln Lys Tyr Gly Glu Ala  
 210 215 220

Lys Ala Gln Glu Leu Phe Ile Asp Asn Pro Arg Lys Ile Val Met Asp  
 225 230 235 240

Gln Leu Ile

&lt;210&gt; 165

&lt;211&gt; 3990

&lt;212&gt; DNA

<213> *Streptococcus pneumoniae*

&lt;400&gt; 165

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3990

<211> 1329

<212> PRT

<213> Streptococcus pneumoniae

<400> 166

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Val Ala Ala Asp Gly Val Thr Pro Thr Thr Thr Glu Asn Gln Pro Thr  
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Ile His Thr Val Ser Asp Ser Pro Gln Ser Ser Glu Asn Arg Thr Glu  
65 70 75 80

Glu Thr Pro Lys Ala Val Leu Gln Pro Glu Ala Pro Lys Thr Val Glu  
85 90 95

Thr Glu Thr Pro Ala Thr Asp Lys Val Ala Ser Leu Pro Lys Thr Glu  
100 105 110

Glu Lys Pro Gln Glu Glu Val Ser Ser Thr Pro Ser Asp Lys Ala Glu  
115 120 125

Val Val Thr Pro Thr Ser Ala Glu Lys Glu Thr Ala Asn Lys Lys Ala  
130 135 140

227

Glu Glu Ala Ser Pro Lys Lys Glu Glu Ala Lys Glu Val Asp Ser Lys  
145 150 155 160

Glu Ser Asn Thr Asp Lys Thr Asp Lys Asp Lys Pro Ala Lys Lys Asp  
165 170 175

Glu Ala Lys Ala Glu Ala Asp Lys Pro Ala Thr Glu Ala Gly Lys Glu  
180 185 190

Arg Ala Ala Thr Val Asn Glu Lys Leu Ala Lys Lys Lys Ile Val Ser  
195 200 205

Ile Asp Ala Gly Arg Lys Tyr Phe Ser Pro Glu Gln Leu Lys Glu Ile  
210 215 220

Ile Asp Lys Ala Lys His Tyr Gly Tyr Thr Asp Leu His Leu Leu Val  
225 230 235 240

Gly Asn Asp Gly Leu Arg Phe Met Leu Asp Asp Met Ser Ile Thr Ala  
245 250 255

Asn Gly Lys Thr Tyr Ala Ser Asp Asp Val Lys Arg Ala Ile Glu Lys  
260 265 270

Gly Thr Asn Asp Tyr Tyr Asn Asp Pro Asn Gly Asn His Leu Thr Glu  
275 280 285

Ser Gln Met Thr Asp Leu Ile Asn Tyr Ala Lys Asp Lys Gly Ile Gly  
290 295 300

Leu Ile Pro Thr Val Asn Ser Pro Gly His Met Asp Ala Ile Leu Asn  
305 310 315 320

Ala Met Lys Glu Leu Gly Ile Gln Asn Pro Asn Phe Ser Tyr Phe Gly  
325 330 335

Lys Lys Ser Ala Arg Thr Val Asp Leu Asp Asn Glu Gln Ala Val Ala  
340 345 350



Phe Thr Lys Ala Leu Ile Asp Lys Tyr Ala Ala Tyr Phe Ala Lys Lys  
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Thr Glu Ile Phe Asn Ile Gly Leu Asp Glu Tyr Ala Asn Asp Ala Thr  
 370 375 380

Asp Ala Lys Gly Trp Ser Val Leu Gln Ala Asp Lys Tyr Tyr Pro Asn  
 385 390 395 400

Glu Gly Tyr Pro Val Lys Gly Tyr Glu Lys Phe Ile Ala Tyr Ala Asn  
 405 410 415

Asp Leu Ala Arg Ile Val Lys Ser His Gly Leu Lys Pro Met Ala Phe  
 420 425 430

Asn Asp Gly Ile Tyr Tyr Asn Ser Asp Thr Ser Phe Gly Ser Phe Asp  
 435 440 445

Lys Asp Ile Ile Val Ser Met Trp Thr Gly Gly Trp Gly Gly Tyr Asp  
 450 455 460

Val Ala Ser Ser Lys Leu Leu Ala Glu Lys Gly His Gln Ile Leu Asn  
 465 470 475 480

Thr Asn Asp Ala Trp Tyr Tyr Val Leu Gly Arg Asn Ala Asp Gly Gln  
 485 490 495

Gly Trp Tyr Asn Leu Asp Gln Gly Leu Asn Gly Ile Lys Asn Thr Pro  
 500 505 510

Ile Thr Ser Val Pro Lys Thr Glu Gly Ala Asp Ile Pro Ile Ile Gly  
 515 520 525

Gly Met Val Ala Ala Trp Ala Asp Thr Pro Ser Ala Arg Tyr Ser Pro  
 530 535 540

Ser Arg Leu Phe Lys Leu Met Arg His Phe Ala Asn Ala Asn Ala Glu  
 545 550 555 560

229

Tyr Phe Ala Ala Asp Tyr Glu Ser Ala Glu Gln Ala Leu Asn Glu Val  
565 570 575

Pro Lys Asp Leu Asn Arg Tyr Thr Ala Glu Ser Val Thr Ala Val Lys  
580 585 590

Glu Ala Glu Lys Ala Ile Arg Ser Leu Asp Ser Asn Leu Ser Arg Ala  
595 600 605

Gln Gln Asp Thr Ile Asp Gln Ala Ile Ala Lys Leu Gln Glu Thr Val  
610 615 620

Asn Asn Leu Thr Leu Thr Pro Glu Ala Gln Lys Glu Glu Glu Ala Lys  
625 630 635 640

Arg Glu Val Glu Lys Leu Ala Lys Asn Lys Val Ile Ser Ile Asp Ala  
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Gly Arg Lys Tyr Phe Thr Leu Asn Gln Leu Lys Arg Ile Val Asp Lys  
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675 680 685

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Ala Tyr Tyr Asp Asp Pro Asn Gly Thr Ala Leu Thr Gln Ala Glu Val  
725 730 735

Thr Glu Leu Ile Glu Tyr Ala Lys Ser Lys Asp Ile Gly Leu Ile Pro  
740 745 750

Ala Ile Asn Ser Pro Gly His Met Asp Ala Met Leu Val Ala Met Glu  
755 760 765

Lys Leu Gly Ile Lys Asn Pro Gln Ala His Phe Asp Lys Val Ser Lys  
 770 775 780

Thr Thr Met Asp Leu Lys Asn Glu Glu Ala Met Asn Phe Val Lys Ala  
 785 790 795 800

Leu Ile Gly Lys Tyr Met Asp Phe Phe Ala Gly Lys Thr Lys Ile Phe  
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Asn Phe Gly Thr Asp Glu Tyr Ala Asn Asp Ala Thr Ser Ala Gln Gly  
 820 825 830

Trp Tyr Tyr Leu Lys Trp Tyr Gln Leu Tyr Gly Lys Phe Ala Glu Tyr  
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Ala Asn Thr Leu Ala Ala Met Ala Lys Glu Arg Gly Leu Gln Pro Met  
 850 855 860

Ala Phe Asn Asp Gly Phe Tyr Tyr Glu Asp Lys Asp Asp Val Gln Phe  
 865 870 875 880

Asp Lys Asp Val Leu Ile Ser Tyr Trp Ser Lys Gly Trp Trp Gly Tyr  
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Asn Leu Ala Ser Pro Gln Tyr Leu Ala Ser Lys Gly Tyr Lys Phe Leu  
 900 905 910

Asn Thr Asn Gly Asp Trp Tyr Tyr Ile Leu Gly Gln Lys Pro Glu Asp  
 915 920 925

Gly Gly Gly Phe Leu Lys Lys Ala Ile Glu Asn Thr Gly Lys Thr Pro  
 930 935 940

Phe Asn Gln Leu Ala Ser Thr Lys Tyr Pro Glu Val Asp Leu Pro Thr  
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Val Gly Ser Met Leu Ser Ile Trp Ala Asp Arg Pro Ser Ala Glu Tyr  
 965 970 975

Lys Glu Glu Glu Ile Phe Glu Leu Met Thr Ala Phe Ala Asp His Asn  
 980 985 990

Lys Asp Tyr Phe Arg Ala Asn Tyr Asn Ala Leu Arg Glu Glu Leu Ala  
 995 1000 1005

Lys Ile Pro Thr Asn Leu Glu Gly Tyr Ser Lys Glu Ser Leu Glu Ala  
 1010 1015 1020

Leu Asp Ala Ala Lys Thr Ala Leu Asn Tyr Asn Leu Asn Arg Asn Lys  
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Gln Ala Glu Leu Asp Thr Leu Val Ala Asn Leu Lys Ala Ala Leu Gln  
 1045 1050 1055

Gly Leu Lys Pro Ala Val Thr His Ser Gly Ser Leu Asp Glu Asn Glu  
 1060 1065 1070

Val Ala Ala Asn Val Glu Thr Arg Pro Glu Leu Ile Thr Arg Thr Glu  
 1075 1080 1085

Glu Ile Pro Phe Glu Val Ile Lys Lys Glu Asn Pro Asn Leu Pro Ala  
 1090 1095 1100

Gly Gln Glu Asn Ile Ile Thr Ala Gly Val Lys Gly Glu Arg Thr His  
 1105 1110 1115 1120

Tyr Ile Ser Val Leu Thr Glu Asn Gly Lys Thr Thr Glu Thr Val Leu  
 1125 1130 1135

Asp Ser Gln Val Thr Lys Glu Val Ile Asn Gln Val Val Glu Val Gly  
 1140 1145 1150

Ala Pro Val Thr His Lys Gly Asp Glu Ser Gly Leu Ala Pro Thr Thr  
 1155 1160 1165

Glu Val Lys Pro Arg Leu Asp Ile Gln Glu Glu Glu Ile Pro Phe Thr  
 1170 1175 1180

232

Thr Val Thr Cys Glu Asn Pro Leu Leu Leu Lys Gly Lys Thr Gln Val  
1185 1190 1195 1200

Ile Thr Lys Gly Val Asn Gly His Arg Ser Asn Phe Tyr Ser Val Ser  
1205 1210 1215

Thr Ser Ala Asp Gly Lys Glu Val Lys Thr Leu Val Asn Ser Val Val  
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Ala Gln Glu Ala Val Thr Gln Ile Val Glu Val Gly Thr Met Val Thr  
1235 1240 1245

His Val Gly Asp Glu Asn Gly Gln Ala Ala Ile Ala Glu Glu Lys Pro  
1250 1255 1260

Lys Leu Glu Ile Pro Ser Gln Pro Ala Pro Ser Thr Ala Pro Ala Glu  
1265 1270 1275 1280

Glu Ser Lys Val Leu Pro Gln Asp Pro Ala Pro Val Val Thr Glu Lys  
1285 1290 1295

Lys Leu Pro Glu Thr Gly Thr His Asp Ser Ala Gly Leu Val Val Ala  
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Gly Leu Met Ser Thr Leu Ala Ala Tyr Gly Leu Thr Lys Arg Lys Glu  
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Asp

<210> 167

<211> 825

<212> DNA

<213> Streptococcus pneumoniae

<400> 167

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acaaagacag tgggcaagga aactgttcca accgtagcta atgcctctctt gtctaaggcc 360
actcgtcagt acaagaatcg taaagaaact gggaatgggt caacttcttg gactcctcca 420
ggttggcatc aggtcaagaa tctaaagggc tcttataccc atgcagtcga tagaggtcac 480
ttgttaggct atgccttaat cgggtggtttg gatgggttttg atgcctcaac aagcaatcct 540
aaaaacattg ctgttcagac agcctgggca aatcaggcac aagccgagta ttcgactgggt 600
caaaactact atgaaagcaa ggtgcgtaaa gccttggacc aaaacaagcg tgtccgttac 660
cgtgtaaccc ttactacgc ttcaaacgag gatttagttc cctcagcttc acagattgaa 720
gccaagtctt cggatggaga attggaattc aatgttctag ttcccaatgt tcaaaagga 780
cttcaactgg attaccgaac tggagaagta actgtaactc agtaa 825

```

<210> 168

<211> 274

<212> PRT

<213> Streptococcus pneumoniae

<400> 168

```

Met Asn Lys Lys Thr Arg Gln Thr Leu Ile Gly Leu Leu Val Leu Leu
  1              5              10              15

```

```

Leu Leu Ser Thr Gly Ser Tyr Tyr Ile Lys Gln Met Pro Ser Ala Pro
      20              25              30

```

```

Asn Ser Pro Lys Thr Asn Leu Ser Gln Lys Lys Gln Ala Ser Glu Ala
      35              40              45

```

```

Pro Ser Gln Ala Leu Ala Glu Ser Val Leu Thr Asp Ala Val Lys Ser
      50              55              60

```

```

Gln Ile Lys Gly Ser Leu Glu Trp Asn Gly Ser Gly Ala Phe Ile Val
      65              70              75              80

```

```

Asn Gly Asn Lys Thr Asn Leu Asp Ala Lys Val Ser Ser Lys Pro Tyr
      85              90              95

```

Ala Asp Asn Lys Thr Lys Thr Val Gly Lys Glu Thr Val Pro Thr Val  
 100 105 110

Ala Asn Ala Leu Leu Ser Lys Ala Thr Arg Gln Tyr Lys Asn Arg Lys  
 115 120 125

Glu Thr Gly Asn Gly Ser Thr Ser Trp Thr Pro Pro Gly Trp His Gln  
 130 135 140

Val Lys Asn Leu Lys Gly Ser Tyr Thr His Ala Val Asp Arg Gly His  
 145 150 155 160

Leu Leu Gly Tyr Ala Leu Ile Gly Gly Leu Asp Gly Phe Asp Ala Ser  
 165 170 175

Thr Ser Asn Pro Lys Asn Ile Ala Val Gln Thr Ala Trp Ala Asn Gln  
 180 185 190

Ala Gln Ala Glu Tyr Ser Thr Gly Gln Asn Tyr Tyr Glu Ser Lys Val  
 195 200 205

Arg Lys Ala Leu Asp Gln Asn Lys Arg Val Arg Tyr Arg Val Thr Leu  
 210 215 220

Tyr Tyr Ala Ser Asn Glu Asp Leu Val Pro Ser Ala Ser Gln Ile Glu  
 225 230 235 240

Ala Lys Ser Ser Asp Gly Glu Leu Glu Phe Asn Val Leu Val Pro Asn  
 245 250 255

Val Gln Lys Gly Leu Gln Leu Asp Tyr Arg Thr Gly Glu Val Thr Val  
 260 265 270

Thr Gln

<210> 169

<211> 225

<212> DNA

<213> Streptococcus pneumoniae

<400> 169

```
gtgctaagat tcagcggatt gaggcaagtg atgaagatga ataagaaatc aagctacgta 60
gtcaagcggt tacttttagt catcatagta ctgatttttag gtactctggc tctaggaatc 120
ggtttaaatg taggttatgg aatcttgggc aagggtcaag atccatgggc tatcctgtct 180
ccagcaaaat ggcaggaatt gattcataaa ttacaggaa attag 225
```

<210> 170

<211> 74

<212> PRT

<213> Streptococcus pneumoniae

<400> 170

```
Val Leu Arg Phe Ser Gly Leu Arg Gln Val Met Lys Met Asn Lys Lys
  1             5             10             15
```

```
Ser Ser Tyr Val Val Lys Arg Leu Leu Leu Val Ile Ile Val Leu Ile
      20             25             30
```

```
Leu Gly Thr Leu Ala Leu Gly Ile Gly Leu Met Val Gly Tyr Gly Ile
      35             40             45
```

```
Leu Gly Lys Gly Gln Asp Pro Trp Ala Ile Leu Ser Pro Ala Lys Trp
      50             55             60
```

```
Gln Glu Leu Ile His Lys Phe Thr Gly Asn
      65             70
```

<210> 171

<211> 40

<212> DNA

<213> Artificial Sequence



<220>

<223> Description of Artificial Sequence: Primer

<400> 171

cgagatctga tatctcacia acagataacg gcgtaaatag

40

<210> 172

<211> 43

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 172

gaagatcttc cccgggatca caaacagata acggcgtaaa tag

43

<210> 173

<211> 42

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 173

cgagatctga tatccatcac aaacagataa cggcgtaaata ag

42

<210> 174

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

&lt;400&gt; 174

cgggatcctt atggacctga atcagcggtg tc

32

&lt;210&gt; 175

&lt;211&gt; 23

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Primer

&lt;400&gt; 175

ggatgctttg tttcaggtgt atc

23

&lt;210&gt; 176

&lt;211&gt; 82

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Primer

&lt;400&gt; 176

catgatatcg gtacctcaag ctcatatcat tgtccggcaa tgggtgtgggc tttttttggt 60

ttagcggata acaatttcac ac

82

&lt;210&gt; 177

&lt;211&gt; 81

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Primer

<400> 177

gcggatcccc cgggcttaat taatgtttaa acactagtcg aagatctcgc gaattctcct 60  
gtgtgaaatt gttatccgct a 81

<210> 178

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 178

cgccagggtt ttcccagtca cgac 24

<210> 179

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 179

tcaggggggc ggagcctatg 20

<210> 180

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 180

tcgtatgttg tgtggaattg tg 22

<210> 181

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 181

tccggctcgt atgttggtg gaattg

26

<210> 182

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<221> SITE

<222> (3)

<223> Xaa=Any amino acid

<220>

<223> Description of Artificial Sequence: Cell wall  
anchoring motif

<400> 182

Leu Pro Xaa Thr Gly

1

5

<210> 183

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 183

gcgggatccg ccaccatg

18

<210> 184

<211> 10

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 184

ttgcggccgc

10

<210> 185

<211> 43

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 185

cggatccgcc accatgggtc taattgaaga cttaaaaaat caa

43

<210> 186

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 186

ttgcggccgc caatgctaga ctaaacacaa gactca

36

<210> 187

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 187

cgcggatcca tgaaaaaaat ctattcattt ttagca

36

<210> 188

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 188

ccctcgaggg ctacttccga tacattttaa actgtagg

38

<210> 189

<211> 35

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 189

cggatccgcc accatgagtc atgtcgctgc aaatg

35

<210> 190

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 190

ttgcggccgc ataccaaacg ctgacatcta cg

32

<210> 191

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 191

cggatccgcc accatgcaaa aagagcggta tggttatg

38

<210> 192

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 192

ttgcggccgc acccccatc ttaatccott

30

<210> 193

<211> 40

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 193

cggatccgcc accatggagg tatgtgaaat gtcacgtaaa

40

<210> 194

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 194

ttgcggccgc ttttaciaaag tcaagcaaag cc

32

<210> 195

<211> 48

<212> PRT

<213> Streptococcus pneumoniae

<400> 195

Gly Ile Arg Leu Arg Asn Met Leu Phe Lys Ile Trp Pro Ala Val Ala

1

5

10

15

Leu Val Thr Ser Ser Gly Asn Asn Val Ser Met Leu His Ser Ile Ala

20

25

30

Asn Met Gly Gln Leu Thr Leu Gly Thr Gln Cys Gln Thr Val Val Val

35

40

45



<210> 196

<211> 11

<212> PRT

<213> Streptococcus pneumoniae

<400> 196

Gln Lys Ile Thr Met Ile Thr Phe Thr Phe Gln

1

5

10